Patent and Trademark Office

## SEARCH REQUEST FORM

Requestor's Name:		rial mber:	
Date:	Phone:	Art Unit:	
hat may have a special meaning. G	search topic. Describe specifically as posive examples or relevant citations, author include a copy of the broadest and/or me	s keywords, etc., if known. Fo	earched. Define any terms or sequences, please attach
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rate completed: 03-28	STAFF USE ONL	yaya asa san	
earcher: Beverly	C-4954 Search Site	Ver STIC	idors IG Suite
erminal time: 24		CM-1	STN
lapsed time: PU time:	Type of Sea	Pre-S	Dialog APS
otal time: 27		N.A. Sequence	Geninfo
Tumber of Searches:	<b>7</b> _	A.A. Sequence —	SDC DARC/Questel ,
Number of Databases:		Structure Bibliographic	Other CGN

PTO-1590 (9-90)

USCOMM-DC 90-3952

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FILE 'REGISTRY' ENTERED AT 14:26:15 ON 28 MAR 2003
            13 S HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSG/SQSP
L1
    ANSWER 1 OF 13 REGISTRY COPYRIGHT 2003 ACS
L1
    474444-80-9 REGISTRY
RN
    L-Serine, L-histidyl-L-seryl-L-.alpha.-aspartylglycyl-L-threonyl-L-
CN
    phenylalanyl-L-threonyl-L-seryl-L-.alpha.-aspartyl-L-leucyl-L-seryl-
    L-lysyl-L-glutaminyl-L-methionyl-L-.alpha.-glutamyl-L-.alpha.-
    glutamyl-L-.alpha.-glutamyl-L-alanyl-L-valyl-L-arginyl-L-leucyl-L-
    phenylalanyl-L-isoleucyl-L-.alpha.-glutamyl-L-tryptophyl-L-leucyl-L-
    lysyl-L-asparaginylglycylglycyl-L-prolyl-L-seryl-L-serylglycyl-L-
    alanyl-L-prolyl-L-prolyl-L-prolyl- (9CI) (CA INDEX NAME)
OTHER NAMES:
    4: PN: WO02085406 SEQID: 7 unclaimed protein
CI
    MAN
SQL
    39
        1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPS
SEO
           HITS AT:
          1 - 34
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
           1: 137:346934
REFERENCE
    ANSWER 2 OF 13 REGISTRY COPYRIGHT 2003 ACS
     437134-20-8 REGISTRY
RN
    8: PN: WO0246227 SEQID: 9 unclaimed protein (9CI) (CA INDEX NAME)
CN
CI
    MAN
    39
SQL
         1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPS
SEO
           _____ =====
          1-34
HITS AT:
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
           1: 137:28591
REFERENCE
     ANSWER 3 OF 13 REGISTRY COPYRIGHT 2003 ACS
T.1
     421042-94-6 REGISTRY
RN
     L-Serine, L-histidyl-L-seryl-L-.alpha.-aspartylglycyl-L-threonyl-L-
CN
     phenylalanyl-L-threonyl-L-seryl-L-.alpha.-aspartyl-L-leucyl-L-seryl-
     L-lysyl-L-glutaminyl-L-methionyl-L-.alpha.-glutamyl-L-.alpha.-
     glutamyl-L-.alpha.-glutamyl-L-alanyl-L-valyl-L-arginyl-L-leucyl-L-
     phenylalanyl-L-isoleucyl-L-.alpha.-glutamyl-L-tryptophyl-L-leucyl-L-
     lysyl-L-asparaginylglycylglycyl-L-prolyl-L-seryl-L-serylglycyl-L-
     alanyl-L-prolyl-L-prolyl- (9CI) (CA INDEX NAME)
     2: PN: WO0234285 SEQID: 7 unclaimed protein
CT
     MAN
SQL
     39
         1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPS
SEO
           HITS AT:
           1 - 34
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
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Searcher: Shears 308-4994

\*REFERENCE 1: 136:350560 ANSWER 4 OF 13 REGISTRY COPYRIGHT 2003 ACS L1309729-82-6 REGISTRY CN Exendin 3 (Heloderma horridum), 39a-[N6-[21-(2,5-dihydro-2,5-dioxo-1H-pyrrol-1-yl)-1,10,19-trioxo-3,6,12,15-tetraoxa-9,18-diazaheneicos-1-yl]-L-lysinamide]-, pentakis(trifluoroacetate) (salt) (9CI) (CA INDEX NAME) OTHER NAMES: CN 38: PN: WO0069911 PAGE: 71 claimed sequence SOL SEO 1 HSDGTFTSDL SKOMEEEAVR LFIEWLKNGG PSSGAPPPSK HITS AT: 1-34 \*\*RELATED SEQUENCES AVAILABLE WITH SEOLINK\*\* SEQ 1 HSDGTFTSDL SKOMEEEAVR LFIEWLKNGG PSSGAPPPSK HITS AT: 1 - 34\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\* SEO 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK HITS AT: 1 - 34\*\*RELATED ŞEQUENCES AVAILABLE WITH SEQLINK\*\* REFERENCE 1: 134:13338 L1ANSWER 5 OF 13 REGISTRY COPYRIGHT 2003 ACS RN 309729-80-4 REGISTRY Exendin 3 (Heloderma horridum), 39a-[N6-[21-(2,5-dihydro-2,5-dioxo-1H-pyrrol-1-yl)-1,10,19-trioxo-3,6,12,15-tetraoxa-9,18-diazaheneicos-1-yl]-L-lysinamide]- (9CI) (CA INDEX NAME) COM, MAN CI SOL 40 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK HITS AT: 1 - 34\*\*RELATED SEQUENCES AVAILABLE WITH SEOLINK\*\* REFERENCE 1: 134:13338 L1ANSWER 6 OF 13 REGISTRY COPYRIGHT 2003 ACS RN 309257-17-8 REGISTRY CN 194: PN: WO0069900 SEQID: 373 unclaimed protein (9CI) (CA INDEX NAME) CI MAN SQL 40

Searcher: Shears 308-4994

1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK

SEO

HITS AT: 1-34

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

REFERENCE 1: 134:21425

L1 ANSWER 7 OF 13 REGISTRY COPYRIGHT 2003 ACS

RN 308806-01-1 REGISTRY

CN L-Lysine, L-histidyl-L-seryl-L-alpha.-aspartylglycyl-L-threonyl-L-phenylalanyl-L-threonyl-L-seryl-L-alpha.-aspartyl-L-leucyl-L-seryl-L-lysyl-L-glutaminyl-L-methionyl-L-alpha.-glutamyl-L-alpha.-glutamyl-L-alpha.-glutamyl-L-leucyl-L-phenylalanyl-L-isoleucyl-L-alpha.-glutamyl-L-tryptophyl-L-leucyl-L-lysyl-L-asparaginylglycylglycyl-L-prolyl-L-seryl-L-serylglycyl-L-alanyl-L-prolyl-L-prolyl-L-seryl- (9CI) (CA INDEX NAME)

OTHER NAMES:

CN 19: PN: WO0069911 SEQID: 19 claimed protein

CI MAN SQL 40

SEO 1 HSDGTFTSDL SKOMEEEAVR LFIEWLKNGG PSSGAPPPSK

HITS AT: 1-34

\*\*RELATED SEOUENCES AVAILABLE WITH SEOLINK\*\*

REFERENCE 1: 134:13338

L1 ANSWER 8 OF 13 REGISTRY COPYRIGHT 2003 ACS

RN 308249-28-7 REGISTRY

CN Exendin 3 (Heloderma horridum), 39a-[N6-[15-(2,5-dihydro-2,5-dioxo-1H-pyrrol-1-yl)-1,7,13-trioxo-3,9-dioxa-6,12-diazapentadec-1-yl]-L-lysinamide]-, pentakis(trifluoroacetate) (salt) (9CI) (CA INDEX NAME)

SQL 40

SEQ 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK

HITS AT: 1-34

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

SEQ 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK

HITS AT: 1-34

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

SEQ 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK

HITS AT: 1-34

\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

REFERENCE 1: 134:21425

L1 ANSWER 9 OF 13 REGISTRY COPYRIGHT 2003 ACS

RN 308249-20-9 REGISTRY

Searcher: Shears 308-4994

Exendin 3 (Heloderma horridum), 39a-[N6-[15-(2,5-dihydro-2,5-dioxo-CN 1H-pyrrol-1-yl)-1,7,13-trioxo-3,9-dioxa-6,12-diazapentadec-1-yl]-Llysinamide] - (9CI) (CA INDEX NAME) COM, MAN CI SOL 40 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK SEO HITS AT: 1-34 \*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\* ANSWER 10 OF 13 REGISTRY COPYRIGHT 2003 ACS 308249-19-6 REGISTRY RN Exendin 3 (Heloderma horridum), 39a-[N6-[3-(2,5-dihydro-2,5-dioxo-1H-CN pyrrol-1-yl)-1-oxopropyl]-L-lysinamide]-, pentakis(trifluoroacetate) (salt) (9CI) (CA INDEX NAME) OTHER NAMES: 35: PN: WO0069911 PAGE: 68 claimed sequence CN SQL 40 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK SEO \_ HITS AT: 1 - 34\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\* 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK SEQ HITS AT: 1 - 34\*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\* 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK SEO \_\_\_\_\_ 1-34 HITS AT: \*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\* REFERENCE 1: 134:21425 REFERENCE 2: 134:13338 ANSWER 11 OF 13 REGISTRY COPYRIGHT 2003 ACS L1RN 308245-55-8 REGISTRY Exendin 3 (Heloderma horridum), 39a-[N6-[3-(2,5-dihydro-2,5-dioxo-1Hpyrrol-1-yl)-1-oxopropyl]-L-lysinamide]- (9CI) (CA INDEX NAME) COM, MAN CI SQL 40 1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPSK SEQ \_ 1-34 HITS AT: \*\*RELATED SEQUENCES AVAILABLE WITH SEQLINK\*\*

Searcher: Shears 308-4994

REFERENCE 1: 134:13338

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ANSWER 12 OF 13 REGISTRY COPYRIGHT 2003 ACS
L1
     306277-48-5 REGISTRY
RN
     L-Serine, L-histidyl-L-seryl-L-.alpha.-aspartylglycyl-L-threonyl-L-
CN
     phenylalanyl-L-threonyl-L-seryl-L-.alpha.-aspartyl-L-leucyl-L-seryl-
     L-lysyl-L-glutaminyl-L-methionyl-L-.alpha.-glutamyl-L-.alpha.-
     glutamyl-L-.alpha.-glutamyl-L-alanyl-L-valyl-L-arginyl-L-leucyl-L-
     phenylalanyl-L-isoleucyl-L-.alpha.-glutamyl-L-tryptophyl-L-leucyl-L-
     lysyl-L-asparaginylglycylglycyl-L-prolyl-L-seryl-L-serylglycyl-L-
     alanyl-L-prolyl-L-prolyl-L-prolyl- (9CI) (CA INDEX NAME)
OTHER NAMES:
CN
     11: PN: WO0069911 SEQID: 11 claimed protein
     185: PN: WO0069900 SEQID: 364 unclaimed protein
CN
     1: PN: WO0151078 SEQID: 1 unclaimed protein
CN
     2: PN: WO0066138 PAGE: 13 unclaimed protein
CN
     3: PN: US6284725 SEQID: 7 unclaimed protein
CN
     3: PN: WO0066142 TABLE: 1 unclaimed protein
CN
     5: PN: WO0077039 TABLE: 1 unclaimed protein
CN
     MAN
CI
SOL
     39
         1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPS
SEO
           ________________________________
HITS AT:
           1 - 34
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
               135:205920
            1:
REFERENCE
REFERENCE
            2:
               135:132445
REFERENCE
            3:
               134:51920
            4:
               134:21425
REFERENCE
REFERENCE
            5:
               134:13338
                133:345167
REFERENCE
            6:
REFERENCE
               133:345166
            7.:
     ANSWER 13 OF 13 REGISTRY COPYRIGHT 2003 ACS
     130357-25-4 REGISTRY
     Exendin 3 (Heloderma horridum) (9CI) (CA INDEX NAME)
OTHER NAMES:
     11: PN: WO0041546 FIGURE: 1 claimed protein
     1: PN: WO0066629 FIGURE: 1 unclaimed sequence
     36: PN: WO0069911 PAGE: 69 claimed sequence
CN
     Exendin-3 (Heloderma horridum)
CN
     L-Serinamide, L-histidyl-L-seryl-L-.alpha.-aspartylglycyl-L-threonyl-
CN
     L-phenylalanyl-L-threonyl-L-seryl-L-.alpha.-aspartyl-L-leucyl-L-
     seryl-L-lysyl-L-glutaminyl-L-methionyl-L-.alpha.-glutamyl-L-.alpha.-
     glutamyl-L-.alpha.-glutamyl-L-alanyl-L-valyl-L-arginyl-L-leucyl-L-
     phenylalanyl-L-isoleucyl-L-.alpha.-glutamyl-L-tryptophyl-L-leucyl-L-
     lysyl-L-asparaginylglycylglycyl-L-prolyl-L-seryl-L-serylglycyl-L-
     alanyl-L-prolyl-L-prolyl-
CI
     MAN
SQL
     39
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Searcher: Shears 308-4994

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1 HSDGTFTSDL SKQMEEEAVR LFIEWLKNGG PSSGAPPPS
SEO
            HITS AT:
            1 - 34
**RELATED SEQUENCES AVAILABLE WITH SEQLINK**
REFERENCE
             1: 137:88643
             2: 135:376777
REFERENCE
             3: 134:37033
REFERENCE
             4: 134:13338
REFERENCE
              5: 133:359242
REFERENCE
REFERENCE
              6: 133:135612
              7: 133:129880
REFERENCE
REFERENCE
              8: 128:192936
REFERENCE
              9: 123:74913
REFERENCE 10: 121:222490
      FILE 'HCAPLUS' ENTERED AT 14:27:42 ON 28 MAR 2003
               20 S L1
L2
     ANSWER 1 OF 20 HCAPLUS COPYRIGHT 2003 ACS
                             2002:832649 HCAPLUS
ACCESSION NUMBER:
                             137:346934
DOCUMENT NUMBER:
                             Methods for treating conditions associated with
TITLE:
                             insulin resistance by administering a GLP-1
                             compound
                             Holst, Jens Juul; Olsen, Mette Zander; Hathaway,
INVENTOR(S):
                             David R.
                             Restoragen, Inc., USA
PATENT ASSIGNEE(S):
                             PCT Int. Appl., 60 pp.
SOURCE:
                             CODEN: PIXXD2
DOCUMENT TYPE:
                             Patent
                             English
LANGUAGE:
FAMILY ACC. NUM. COUNT: 1
PATENT INFORMATION:
                                                APPLICATION NO. DATE
      PATENT NO.
                     KIND DATE
                                                 _____
                                _____
      _____ ___
                                                WO 2002-US13088 20020424
                                 20021031
      WO 2002085406
                         A1
               AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH,
               CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ,
               NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
          RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT,
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Searcher: Shears 308-4994

SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE,

```
SN, TD, TG
                                          US 2001-285699P P 20010424
PRIORITY APPLN. INFO.:
    The present invention relates to methods and compns. for treating
     insulin-assocd. conditions comprising administering a glucagon-like
     peptide-1 (GLP-1) compd. to subjects suffering therefrom. The
     insulin resistance-assocd. condition of the invention is type-2
     pre-diabetes, atherosclerotic cardiovascular disease, drug-induced
     insulin resistance, congestive heart failure, diminished exercise
     capacity of skeletal muscle, and left ventricular dysfunction with
     cardiac metabolic myopathy or diminished exercise capacity of
     skeletal muscle; with the proviso that said congestive heart failure
     is not assocd. with toxic hypervolemia.
     474444-80-9
IT
     RL: PRP (Properties)
        (unclaimed protein sequence; methods for treating conditions
        assocd. with insulin resistance by administering a GLP-1 compd.)
                                 THERE ARE 11 CITED REFERENCES AVAILABLE
                          11
REFERENCE COUNT:
                                 FOR THIS RECORD. ALL CITATIONS AVAILABLE
                                 IN THE RE FORMAT
     ANSWER 2 OF 20 HCAPLUS COPYRIGHT 2003 ACS
                          2002:449715 HCAPLUS
ACCESSION NUMBER:
                          137:28591
DOCUMENT NUMBER:
                          Preparation of GLP-1 fusion proteins for use in
TITLE:
                          treating diabetes mellitus and other conditions
                          Glaesner, Wolfgang; Micanovic, Radmilla;
INVENTOR(S):
                          Tschang, Sheng-Hung Rainbow
                          Eli Lilly and Company, USA
PATENT ASSIGNEE(S):
                          PCT Int. Appl., 200 pp.
SOURCE:
                          CODEN: PIXXD2
                          Patent
DOCUMENT TYPE:
                          English
LANGUAGE:
FAMILY ACC. NUM. COUNT: 1
PATENT INFORMATION:
                                            APPLICATION NO. DATE
                     KIND DATE
     PATENT NO.
     _____ ___
                                             _____
                                            WO 2001-US43165 20011129
     WO 2002046227
                      A2 20020613
         W: AE, AG, AL, AM, AT, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA,
             CH, CN, CO, CR, CU, CZ, CZ, DE, DK, DK, DM, DZ, EC, EE, EE, ES, FI, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG,
              MK, MN, MW, MX, MZ, NO, NZ, PH, PL, PT, RO, RU, SD, SE, SG,
              SI, SK, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU,
              ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ
         RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT,
              SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE,
              SN, TD, TG
                                                                20011129
                                              AU 2002-26897
                              20020618
     AU 2002026897
                       Α5
                                           US 2000-251954P P
                                                                20001207
PRIORITY APPLN. INFO.:
                                           WO 2001-US43165 W 20011129
                          MARPAT 137:28591
OTHER SOURCE(S):
     The present invention relates to glucagon-like peptide-1 compds.
      fused to proteins that have the effect of extending the in vivo
     half-life of the peptides. The heterologous fusion proteins of the
      invention comprise a GLP-1 compd. fused to human albumin, a human
```

Searcher: Shears 308-4994

albumin analog or fragment, the Fc portion of an Ig, or an analog or

fragment of the Fc portion of an Ig. These fusion proteins can be used to treat non-insulin dependent diabetes mellitus as well as a variety of other conditions. Pharmaceutical formulations contg. the fusion proteins and polynucleotides encoding the proteins are also claimed.

ΙT 437134-20-8

RL: PRP (Properties)

(unclaimed protein sequence; prepn. of GLP-1 fusion proteins for use in treating diabetes mellitus and other conditions)

ANSWER 3 OF 20 HCAPLUS COPYRIGHT 2003 ACS 2002:361520 HCAPLUS ACCESSION NUMBER:

137:88643 DOCUMENT NUMBER:

TITLE:

Endoproteolysis by isolated membrane peptidases reveal metabolic stability of glucagon-like

peptide-1 analogs, exendins-3 and -4

Thum, A.; Hupe-Sodmann, K.; Goke, R.; Voigt, K.; AUTHOR(S):

Goke, B.; McGregor, G. P.

Institute of Physiology, Philipps-University, CORPORATE SOURCE:

Marburg, D-35037, Germany

Experimental and Clinical Endocrinology & SOURCE:

Diabetes (2002), 110(3), 113-118 CODEN: ECEDFQ; ISSN: 0947-7349

Johann Ambrosius Barth PUBLISHER:

Journal DOCUMENT TYPE: English LANGUAGE:

These in vitro studies aimed to characterize the pattern and the AΒ kinetics of endoproteolysis of the insulinotropic hormone glucagon-like peptide-1 (GLP-1) and related peptides by native ectopeptidases. Peptides were incubated with isolated rat or pig kidney brush-border microvilli membranes, which are a rich source of the ectopeptidases that are responsible for the post-secretory metab. of peptide hormones. The proteolytic products were sepd. by reversed-phase HPLC column chromatog. and characterized by mol. mass and primary structure. The relative importance of specific peptidases was established by measuring the effects of specific peptidase inhibitors on the kinetics of proteolysis. Dipeptidyl-peptidase-IV was found to be rate-limiting in the endoproteolysis of GLP-1. GLP-1 homologs, exendins-3 and -4, exhibited exceptional stability in the presence of isolated kidney microvilli membranes. Our finding that exendin-4 is several orders of magnitude more stable than GLP-1 and Ser-8-GLP-1 is esp. noteworthy given this peptide's widely reported insulinotropic potency.

130357-25-4, Exendin 3 (Heloderma horridum)

RL: BSU (Biological study, unclassified); PKT (Pharmacokinetics); BIOL (Biological study)

(endoproteolysis by isolated membrane peptidases reveal metabolic stability of glucagon-like peptide-1 analogs and exendin-3 and -4)

REFERENCE COUNT:

32 THERE ARE 32 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 4 OF 20 HCAPLUS COPYRIGHT 2003 ACS 2002:332051 HCAPLUS ACCESSION NUMBER:

DOCUMENT NUMBER: 136:350560

Treatment of hibernating myocardium and diabetic TITLE:

> 308-4994 Searcher : Shears

cardiomyopathy with a GLP-1 peptide

INVENTOR(S):

Ehlers, Mario
Coolidge, Thomas R., USA
PCT Int. Appl., 26 pp.
CODEN: PIXXD2 PATENT ASSIGNEE(S): SOURCE:

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

	PATENT	NO.		KII	ND	DATE			Al	PPLI	CATI	ON NO	0.	DATE		
	WO 2002 W:	0342 AE, CN, GE, LC, NO, TR,	85 AG, CO, GH, LK, NZ, TT,	AL, CR, GM, LR, PH, TZ,	AM, CU, HR, LS, PL, UA,	2002 AT, CZ, HU, LT, PT, UG,	0502 AU, DE, ID, LU, RO,	AZ, DK, IL, LV, RU,	BA, DM, IN, MA, SD,	BB, DZ, IS, MD, SE,	BG, EC, JP, MG, SG,	BR, EE, KE, MK, SI,	BY, ES, KG, MN, SK,	2001 BZ, FI, KP, MW, SL, AZ,	CA, GB, KR, MX, TJ,	GD, KZ, MZ, TM,
	RW:	GH, CY,	DE, BF,	KE, DK,	LS, ES,	MW, FI,	FR,	GB,	GR,	ΙE,	IT,	LU,	MC,	AT, NL, MR,	PT,	SE,
PRIO	AU 2002 US 2002 RITY APP	1464	05	A		2002 2002			US 20 US 20 US 20 US 20	S 200 000-2 000-2	01-9 2418 2421 2452	39P 34P	8 P P P	2001 2001 2000 2000 2000 2001	1022 1020 1023 1103	
AB	Hiberna impaire myocyte require such as from hi catecho GLP-1 ra metho cardiom	d fur s rements nore bernalamin educe d of	nctions of epine atinomes in the estimation of t	on decellong consequences of the consequences	ue t lula trac ine, ocar to a no	o loor intion are dium card; repired	cali tegr . H. bel . L. iomy neph:	cter zed ity, igh ieve ikew opat rine	ized reduct but plass d to ise, hy in	by cannot be particularly be particularly be particularly be particularly by the particular by	viab perf not evel pred h le tien and	le musionsust of sust	yoca n. ain cat ve c of ith thus	Hibe high echo f mod diab	m wirnat: -ene: lamin rtal:	ing rgy nes, ity
IT		(Pr	ed p	rote:	in s	eque:								ıg my	ocar	dium
DOCUL TITL INVE PATE SOUR DOCUL LANG FAMI	NTOR(S): NT ASSIG	MBER: BER: NEE(	: S): COUI		135 Pep Hol Bio PCT COD Pat	CO: 1:85 :376 tide mqui nebra Int EN: ent lish	0956 777 pha st, aska . App	HC rmac Bart , In pl.,	eution; I	S cal : Dorma USA						

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APPLICATION NO. DATE
                          KIND DATE
      PATENT NO.
                                                       _____
                           ____
      _____
      WO 2001087322 A2
WO 2001087322 A3
                                                       WO 2001-US15872 20010517
                                    20011122
                           A3 20020718
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                MD, RU, TJ, TM
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                TR, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD,
                                                    US 2001-858880
                          A1 20020523
                                                                              20010517
      US 2002061838
                                                    US 2000-205377P P 20000517
PRIORITY APPLN. INFO.:
                                                    US 2000-205262P P 20000519
      A pharmaceutical compn. for administration to a mammal is disclosed.
AΒ
      The compn. includes a therapeutically effective amt. of a peptide,
      such as a GLP-1 mol., a PTH mol., or a GRF mol. The compn. further
      includes a buffer including a weak acid having an acid dissocn.
      const. value of greater than about 1x10-5, such as acetic acid.
      compn. also includes an excipient for making the compn. generally
      isotonic, such as D-mannitol.
      130357-25-4, Exendin 3 (Heloderma horridum)
TT
      RL: PEP (Physical, engineering or chemical process); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); PROC
       (Process); USES (Uses)
           (peptide pharmaceutical formulations relating to parathormone,
          glucagon-like peptide-1, and growth hormone-releasing factor)
      ANSWER 6 OF 20 HCAPLUS COPYRIGHT 2003 ACS
                                2001:650487 HCAPLUS
ACCESSION NUMBER:
                                135:205920
DOCUMENT NUMBER:
                                Metabolic intervention with GLP-1 to improve the
TITLE:
                                function of ischemic and reperfused tissue
                                Coolidge, Thomas R.; Ehlers, Mario R. W.
INVENTOR(S):
                                BioNebraska, Inc., USA
PATENT ASSIGNEE(S):
                                U.S., 10 pp.
SOURCE:
                                CODEN: USXXAM
DOCUMENT TYPE:
                                Patent
                                English
LANGUAGE:
FAMILY ACC. NUM. COUNT: 3
PATENT INFORMATION:
                                                      APPLICATION NO. DATE
                       KIND DATE
       PATENT NO.
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                                    20010904
                                                      US 1999-302596
                                                                              19990430
       US 6284725 B1
                                                       WO 2000-US11251 20000427
                            A2
                                    20001109
      WO 2000066138
                          A3 20010705
       WO 2000066138
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            CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM

RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,
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Searcher: Shears 308-4994

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DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
                                              EP 2000-926404 20000427
                         A2 20020123
     EP 1173197
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     NZ 514610
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                                              NZ 2000-514610
                                                                 20000427
     JP 2002543142
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                              20021217
                                               JP 2000-615022
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                                              US 2001-851738
     US 2002055460
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                         Α1
                                                                 20010509
                                              US 2001-953021
     US 2002147131
                              20021010
                         Α1
                                                                 20010911
     NO 2001005294
                              20011228
                                              NO 2001-5294
                         Α
                                                                 20011029
                                           US 1998-103498P
                                                             P 19981008
PRIORITY APPLN. INFO.:
                                                              A 19990430
                                           US 1999-302596
                                           WO 2000-US11251 W 20000427
                                           US 2001-851738
                                                             A1 20010509
     Individuals in need of treatment of ischemia-related reperfusion are
AB
     treated, preferably i.v., with a compn. which includes a compd.
     which binds to a receptor for the glucagon-like peptide-1. The
     invention relates to both the method and compns. for such treatment.
ΙT
     306277-48-5
     RL: PRP (Properties)
         (unclaimed protein sequence; metabolic intervention with GLP-1 to
        improve the function of ischemic and reperfused tissue)
REFERENCE COUNT:
                                  THERE ARE 7 CITED REFERENCES AVAILABLE FOR
                                  THIS RECORD. ALL CITATIONS AVAILABLE IN
                                  THE RE FORMAT
     ANSWER 7 OF 20 HCAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER:
                           2001:525943 HCAPLUS
DOCUMENT NUMBER:
                           135:132445
TITLE:
                           Use of exendins and agonists thereof for
                           modulation of triglyceride levels and treatment
                           of dyslipidemia
                           Kolterman, Orville Gene; Young, Andrew A.
INVENTOR(S):
                           Amylin Pharmaceuticals, Inc., USA
PATENT ASSIGNEE(S):
                           PCT Int. Appl., 161 pp.
SOURCE:
                           CODEN: PIXXD2
DOCUMENT TYPE:
                           Patent
LANGUAGE:
                           English
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
                                             APPLICATION NO. DATE
     PATENT NO.
                      KIND DATE
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                      A1 20010719
                                             WO 2001-US719 20010109
     WO 2001051078
         W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH,
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              GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK,
              LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ,
              PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ,
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R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO, MK, CY, AL, TR

EP 2001-900978

EP 1246638

A1

20021009

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20030220
                                           US 2001-756690
                                                             20010109
     US 2003036504
                       Α1
                                         US 2000-175365P P 20000110
PRIORITY APPLN. INFO.:
                                                         W 20010109
                                         WO 2001-US719
     Methods for modulating the levels of plasma triglyceride and other
AB
     lipids in a subject comprise administration of an effective amt. of
     an exendin or exendin agonist, alone or in conjunction with other
     compds. or compns. that lower blood triglyceride and/or other lipid
     levels.
     306277-48-5
TΤ
     RL: PRP (Properties)
        (unclaimed protein sequence; use of exendins and agonists thereof
        for modulation of triglyceride levels and treatment of
        dyslipidemia)
                               THERE ARE 7 CITED REFERENCES AVAILABLE FOR
REFERENCE COUNT:
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                               THE RE FORMAT
     ANSWER 8 OF 20 HCAPLUS COPYRIGHT 2003 ACS
L2
                         2000:900675 HCAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                         134:51920
                         GLP-1 as a diagnostic test to determine
TITLE:
                          .beta.-cell function and the presence of
                         impaired glucose tolerance (IGT) and type-II
                         diabetes
                         Holst, J. J.; Vilsboll, Tina
INVENTOR(S):
                         Bionebraska, Inc., USA
PATENT ASSIGNEE(S):
                         PCT Int. Appl., 42 pp.
SOURCE:
                         CODEN: PIXXD2
                         Patent
DOCUMENT TYPE:
                         English
LANGUAGE:
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
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                                            WO 2000-US16428 20000614
     WO 2000077039
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             LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU,
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              PT, IE, SI, LT, LV, FI, RO
                                         US 1999-333415 A 19990615
WO 2000-US16428 W 20000614
PRIORITY APPLN. INFO.:
     Since glucagon-like peptide-1 (GLP-1) is the most potent
AB
     insulinotropic hormone known and has been shown to stimulate insulin
     secretion strongly in patients with type II diabetes, this invention
     uses GLP-1 or its biol. active analogs in .beta.-cell stimulatory
     tests in order to test .beta.-cell function in a simple way. The
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Searcher: Shears 308-4994

test provides information about insulin secretory capacity, is easy

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and reproducible and has insignificant side effects.
     306277-48-5
IT
     RL: PRP (Properties)
        (unclaimed protein sequence; gLP-1 as a diagnostic test to det.
        .beta.-cell function and the presence of impaired glucose
        tolerance (IGT) and type-II diabetes)
     ANSWER 9 OF 20 HCAPLUS COPYRIGHT 2003 ACS
1.2
ACCESSION NUMBER:
                         2000:861704 HCAPLUS
                         134:37033
DOCUMENT NUMBER:
                         Use of exendins and agonists thereof for the
TITLE:
                         treatment of gestational diabetes mellitus
                         Hiles, Richard; Prickett, Kathryn S.
INVENTOR(S):
                         Amylin Pharmaceuticals, Inc., USA
PATENT ASSIGNEE(S):
                         PCT Int. Appl., 133 pp.
SOURCE:
                         CODEN: PIXXD2
DOCUMENT TYPE:
                         Patent
                         English
LANGUAGE:
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
                                          APPLICATION NO. DATE
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                                           WO 2000-US14231 20000523
                       A2
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             LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL,
             PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA,
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         RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH,
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                                           US 1999-323867
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                                           JP 2001-500655
     JP 2003501361
                       T2
                            20030114
                                        US 1999-323867 A 19990601
PRIORITY APPLN. INFO.:
                                        WO 2000-US14231 W 20000523
     Methods for treating gestational diabetes which comprise
AB
     administration of an effective amt. of an exendin or an exendin
     agonist, alone or in conjunction with other compds. or compns. that
     lower blood glucose levels.
     130357-25-4, Exendin 3 (Heloderma horridum)
ΙT
     RL: BAC (Biological activity or effector, except adverse); BSU
     (Biological study, unclassified); THU (Therapeutic use); BIOL
     (Biological study); USES (Uses)
        (use of exendins and agonists thereof for treatment of
        gestational diabetes mellitus in relation to combination with
        insulin or amylin agonist)
     ANSWER 10 OF 20 HCAPLUS COPYRIGHT 2003 ACS
                         2000:824301 HCAPLUS
ACCESSION NUMBER:
                         134:13338
DOCUMENT NUMBER:
                         Long lasting insulinotropic peptides
TITLE:
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Searcher: Shears 308-4994

INVENTOR(S):

Bridon, Dominique P.; L'Archeveque, Benoit;

Ezrin, Alan M.; Holmes, Darren L.; Leblanc,

Anouk; St. Pierre, Serge

PATENT ASSIGNEE(S):

Conjuchem, Inc., Can. PCT Int. Appl., 96 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

SOURCE:

AB

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

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APPLICATION NO.
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             SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ,
             VN, YU, ZA, ZW
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PRIORITY APPLN. INFO.:
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                                                         A3 20000517
                                        EP 2000-932570
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                                        WO 2000-IB763
                                                         W
                                                         W
                                                            20000517
                                        WO 2000-US13563
                                                         A3 20000905
                                        US 2000-623618
     Modified insulinotropic peptides are disclosed. The modified
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insulinotropic peptides are capable of forming a peptidase stabilized insulinotropic peptide. The modified insulinotropic peptides are capable of forming covalent bonds with one or more blood components to form a conjugate. The conjugates may be formed in vivo or ex vivo. The modified peptides are administered to treat humans with diabetes and other related diseases.

306277-48-5 308245-55-8 308806-01-1 IΤ 309729-80-4

> RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(long lasting insulinotropic peptides with antidiabetic activity)

130357-25-4P, Exendin 3 (Heloderma horridum) ΤT

308249-19-6P 309729-82-6P

RL: SPN (Synthetic preparation); PREP (Preparation)

(long lasting insulinotropic peptides with antidiabetic activity) REFERENCE COUNT: THERE ARE 6 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 11 OF 20 HCAPLUS COPYRIGHT 2003 ACS

ACCESSION NUMBER:

2000:824291 HCAPLUS

DOCUMENT NUMBER:

134:21425

TITLE:

Protection of endogenous therapeutic peptides from peptidase activity through conjugation to

blood components

INVENTOR(S):

Bridon, Dominique P.; Ezrin, Alan M.; Milner, Peter G.; Holmes, Darren L.; Thibaudeau, Karen

PATENT ASSIGNEE(S):

SOURCE:

Conjuchem, Inc., Can. PCT Int. Appl., 733 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

LANGUAGE:

Patent

English FAMILY ACC. NUM. COUNT: 3

PATENT INFORMATION:

PAT	rent	NO.		KI	ND	DATE	DATE APPLICATION NO.									DATE						
WO	2000 2000 2000	0699	00	A: A: C:	3	2000 2001 2002	0215		W	0 20	00-U	s135	76	2000	0517							
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									WO 2000-IB763 20000517													
WO	2000 W:	AE, CU, ID, LU, SD,	AL, CZ, IL, LV,	AM, DE, IN, MA, SG,	AT, DK, IS, MD, SI,	AU, DM, JP, MG, SK,	AZ, EE, KE, MK,	ES, KG, MN,	FI, KP, MW,	GB, KR, MX,	GD, KZ, NO,	GE, LC, NZ,	GH, LK, PL,	GM, LR, PT,	HR, LS, RO,	HU, LT, RU,						

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PRIORITY APPLN. INFO.:
                                        US 1999-153406P P
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                                        US 1999-159783P
                                                        P 19991015
                                                         A3 20000517
                                        EP 2000-932570
                                        WO 2000-IB763
                                                         W
                                                            20000517
                                        WO 2000-US13576 W 20000517
    A method for protecting a peptide from peptidase activity in vivo,
AΒ
    the peptide being composed of between 2 and 50 amino acids and
    having a C-terminus and an N-terminus and a C-terminus amino acid
     and an N-terminus amino acid is described. In the first step of the
    method, the peptide is modified by attaching a reactive group to the
    C-terminus amino acid, to the N-terminus amino acid, or to an amino
     acid located between the N-terminus and the C-terminus, such that
    the modified peptide is capable of forming a covalent bond in vivo
    with a reactive functionality on a blood component. The solid phase
    peptide synthesis of a no. of derivs. with 3-maleimidopropionic acid
     (3-MPA) is described. In the next step, a covalent bond is formed
     between the reactive group and a reactive functionality on a blood
     component to form a peptide-blood component conjugate, thereby
    protecting said peptide from peptidase activity. The final step of
     the method involves the analyzing of the stability of the
    peptide-blood component conjugate to assess the protection of the
    peptide from peptidase activity. Thus, the percentage of a K5
     kringle peptide (Pro-Arg-Lys-Leu-Tyr-Asp-Lys-NH2) conjugated to
     human serum albumin via MPA remained relatively const. through a
     24-h plasma assay in contrast to unmodified K5 which decreased to 9%
     of the original amt. of K5 in only 4 h in plasma.
ΙT
     308249-19-6P 308249-28-7P
     RL: RCT (Reactant); SPN (Synthetic preparation); THU (Therapeutic
     use); BIOL (Biological study); PREP (Preparation); RACT (Reactant or
     reagent); USES (Uses)
        (protection of endogenous therapeutic peptides from peptidase
        activity through conjugation to blood components)
     306277-48-5 309257-17-8
ΙT
     RL: PRP (Properties)
        (unclaimed protein sequence; protection of endogenous therapeutic
        peptides from peptidase activity through conjugation to blood
        components)
```

Searcher: Shears 308-4994

HCAPLUS COPYRIGHT 2003 ACS 2000:790546 HCAPLUS

133:359242

ANSWER 12 OF 20

ACCESSION NUMBER: DOCUMENT NUMBER:

09/756690 Modified exendins and exendin agonists TITLE: Young, Andrew; Prickett, Kathryn INVENTOR(S): Amylin Pharmaceuticals, Inc., USA PATENT ASSIGNEE(S): PCT Int. Appl., 119 pp. SOURCE: CODEN: PIXXD2 DOCUMENT TYPE: Patent English LANGUAGE: FAMILY ACC. NUM. COUNT: 1 PATENT INFORMATION: APPLICATION NO. DATE KIND DATE PATENT NO. ----\_\_\_\_\_ 20001109 WO 2000-US11814 20000428 WO 2000066629 A1 W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG EP 2000-928685 20000428 A1 20020130 EP 1175443 AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO BR 2000-10705 20000428 BR 2000010705 A 20020205 JP 2000-615657 20000428 JP 2002544127 T2 20021224 US 1999-132018P P 19990430 PRIORITY APPLN. INFO.: WO 2000-US11814 W 20000428 Novel modified exendins and exendin agonists having an exendin or AB exendin agonists, compns. and methods are useful in treating

exendin agonist linked to one or more polyethylene glycol polymers, for example, and related formulations and dosages and methods of administration thereof are provided. These modified exendins and diabetes and conditions that would be benefited by lowering plasma glucose or delaying and/or slowing gastric emptying or inhibiting food intake.

130357-25-4, Exendin 3 (Heloderma horridum)

RL: PRP (Properties)

(unclaimed sequence; modified exendins and exendin agonists) THERE ARE 6 CITED REFERENCES AVAILABLE FOR REFERENCE COUNT: 6 THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 13 OF 20 HCAPLUS COPYRIGHT 2003 ACS 2000:790326 HCAPLUS ACCESSION NUMBER:

DOCUMENT NUMBER: 133:345167

Metabolic intervention with GLP-1 or its TITLE:

biologically active analogues to improve the function of the ischemic and reperfused brain

Coolidge, Thomas R.; Ehlers, Mario R. W. INVENTOR(S):

PATENT ASSIGNEE(S): Bionebraska, Inc., USA SOURCE: PCT Int. Appl., 19 pp.

CODEN: PIXXD2

Patent DOCUMENT TYPE: English LANGUAGE:

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

308-4994 Shears Searcher :

```
APPLICATION NO.
                                                                DATE
                       KIND DATE
     PATENT NO.
                       ____
                              _____
                                             WO 2000-US11652 20000501
     WO 2000066142
                        A2
                              20001109
     WO 2000066142
                       A3
                             20020124
         W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR,
             CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU,
             ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT,
             LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU,
             SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN,
             YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
         RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,
              DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF,
              BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
                                            US 1999-303016 19990430
EP 2000-928616 20000501
                       B1 20020806
     US 6429197
     EP 1187628
                       A2
                             20020320
         R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC,
             PT, IE, SI, LT, LV, FI, RO
     JP 2002543145
                       T2 20021217
                                              JP 2000-615026
                                                                20000501
                                             NO 2001-5298
     NO 2001005298
                        Α
                              20011228
                                                                20011029
                                           US 1999-303016
                                                            A 19990430
PRIORITY APPLN. INFO.:
                                         US 1998-103498P P 19981008
                                           WO 2000-US11652 W 20000501
     It has now been discovered that GLP-1 treatment after acute stroke
AΒ
     or hemorrhage, preferably i.v. administration, can be an ideal
     treatment because it provides a means for optimizing insulin
     secretion, increasing brain anabolism, enhancing insulin
     effectiveness by suppressing glucagon, and maintaining euglycemia or
     mild hypoglycemia with no risk of severe hypoglycemia.
ΙT
     306277-48-5
     RL: PRP (Properties)
         (unclaimed protein sequence; metabolic intervention with GLP-1 or
        its biol. active analogs to improve the function of the ischemic
        and reperfused brain)
     ANSWER 14 OF 20 HCAPLUS COPYRIGHT 2003 ACS
                          2000:790323 HCAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                           133:345166
                          Metabolic intervention with GLP-1 to improve the
TITLE:
                           function of ischemic and reperfused tissue
                           Coolidge, Thomas R.; Ehlers, Mario R. W.
INVENTOR(S):
                           Bionebraska, Inc., USA
PATENT ASSIGNEE(S):
                           PCT Int. Appl., 22 pp.
SOURCE:
                           CODEN: PIXXD2
DOCUMENT TYPE:
                           Patent
                           English
LANGUAGE:
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
                     KIND DATE
                                             APPLICATION NO. DATE
     PATENT NO.
                                              -----
                     A2
                                             WO 2000-US11251 20000427
                              20001109
     WO 2000066138
                             20010705
     WO 2000066138
                       АЗ
            AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU,
              SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN,
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Searcher: Shears 308-4994

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YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
         RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
                                            US 1999-302596
     US 6284725
                        В1
                              20010904
                                                                19990430
                                             EP 2000-926404
     EP 1173197
                        A2
                              20020123
                                                                20000427
             AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO
                              20020927
                                              NZ 2000-514610
                                                                20000427
                        Α
     JP 2002543142
                        T2
                              20021217
                                              JP 2000-615022
                                                                20000427
     NO 2001005294
                              20011228
                                              NO 2001-5294
                                                                20011029
                        Α
                                           US 1999-302596
PRIORITY APPLN. INFO.:
                                                             Α
                                                                19990430
                                           US 1998-103498P
                                                            Ρ
                                                                19981008
                                           WO 2000-US11251 W 20000427
     Individuals in need of treatment of ischemia-related reperfusion are
AΒ
     treated, preferably i.v., with a compn. which includes a compd.
     which binds to a receptor for the glucagon-like peptide-1. The
     invention relates to both the method and compns. for such treatment.
TΤ
     306277-48-5
     RL: PRP (Properties)
         (unclaimed sequence; metabolic intervention with GLP-1 to improve
        the function of ischemic and reperfused tissue)
     ANSWER 15 OF 20 HCAPLUS COPYRIGHT 2003 ACS
                          2000:493318 HCAPLUS
ACCESSION NUMBER:
DOCUMENT NUMBER:
                          133:129880
TITLE:
                          Methods using an exendin or related substance
                          for glucagon suppression
INVENTOR(S):
                          Young, Andrew; Gedulin, Bronislava
                          Amylin Pharmaceuticals, Inc., USA
PATENT ASSIGNEE(S):
                          PCT Int. Appl., 96 pp.
SOURCE:
                          CODEN: PIXXD2
DOCUMENT TYPE:
                          Patent
                          English
LANGUAGE:
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
     PATENT NO.
                       KIND
                              DATE
                                             APPLICATION NO.
                                                                DATE
     WO 2000041548
                        A2
                              20000720
                                             WO 2000-US942
                                                                20000114
     WO 2000041548
                        A3
                              20001130
             AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR,
             CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU,
             ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT,
             LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU,
             SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ,
             VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
         RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,
             DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF,
              BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
     CA 2356331
                              20000720
                                            CA 2000-2356331 20000114
                        AA
     EP 1143989
                                             EP 2000-902415
                        A2
                              20011017
                                                                20000114
                              20020911
     EP 1143989
                        А3
             AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC,
             PT, IE, SI, LT, LV, FI, RO
     BR 2000007823
                              20011120
                                              BR 2000-7823
                                                                20000114
                        А
     JP 2002538084
                        Т2
                              20021112
                                              JP 2000-593169
                                                                20000114
     NO 2001003469
                        Α
                              20010914
                                              NO 2001-3469
                                                                20010712
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Searcher: Shears 308-4994

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PRIORITY APPLN. INFO.:
                                         US 1999-116380P P
                                                              19990114
                                         US 1999-132017P P
                                                              19990430
                                         US 2000-175365P P
                                                              20000110
                                                          W 20000114
                                         WO 2000-US942
     Methods are provided for use of an exendin, an exendin agonist, or a
AB
     modified exendin or exendin agonist having an exendin or exendin
     agonist linked to one or more polyethylene glycol polymers, for
     example, for lowering glucagon levels and/or suppressing glucagon
     secretion in a subject. These methods are useful in treating
     hyperglucagonemia and other conditions that would be benefited by
     lowering plasma glucagon or suppressing glucagon secretion.
     130357-25-4P, Exendin 3 (Heloderma horridum)
ΙT
     RL: BAC (Biological activity or effector, except adverse); BSU
     (Biological study, unclassified); PRP (Properties); SPN (Synthetic preparation); THU (Therapeutic use); BIOL (Biological study); PREP
     (Preparation); USES (Uses)
        (exendin or related substance for glucagon suppression)
     ANSWER 16 OF 20 HCAPLUS COPYRIGHT 2003 ACS
                         2000:493315 HCAPLUS
ACCESSION NUMBER:
                         133:135612
DOCUMENT NUMBER:
                         Novel exendin agonist formulations and methods
TITLE:
                         of administration thereof
                         Young, Andrew; L'Italien, James J.; Kolterman,
INVENTOR(S):
                         Orville
                         Amylin Pharmaceuticals, Inc., USA
PATENT ASSIGNEE(S):
                         PCT Int. Appl., 281 pp.
SOURCE:
                         CODEN: PIXXD2
DOCUMENT TYPE:
                         Patent
                         English
LANGUAGE:
FAMILY ACC. NUM. COUNT:
PATENT INFORMATION:
                                          APPLICATION NO. DATE
     PATENT NO.
                     KIND DATE
                     A2 20000720
                                          WO 2000-US902 20000114
     WO 2000041546
         W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR,
             CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU,
             ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT,
             LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU,
             SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ,
             VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
         RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,
             DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF,
             BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
                                            CA 2000-2356706 20000114
                             20000720
     CA 2356706
                       AA
                                            EP 2000-914425
                                                              20000114
                       Α2
                             20011010
     EP 1140145
             AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC,
             PT, IE, SI, LT, LV, FI, RO
                                            BR 2000-7820
                             20011120
                                                              20000114
     BR 2000007820
                      Α
                                            JP 2000-593167
                                                              20000114
                       T2
                             20021015
     JP 2002534450
                                            NO 2001-3468
                                                              20010712
     NO 2001003468
                             20010914
                       Α
                                         US 1999-116380P P
                                                              19990114
PRIORITY APPLN. INFO.:
                                         US 2000-175365P P
                                                              20000110
                                                          W 20000114
                                         WO 2000-US902
     Novel exendin and exendin agonist compd. formulations and dosages
AB
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Searcher: Shears 308-4994

and methods of administration thereof are provided. These compns. and methods are useful in treating diabetes and conditions that

would be benefited by lowering plasma glucose or delaying and/or slowing gastric emptying or inhibiting food intake. 130357-25-4P, Exendin-3 (Heloderma horridum) IT RL: BAC (Biological activity or effector, except adverse); BOC (Biological occurrence); BSU (Biological study, unclassified); PNU (Preparation, unclassified); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); OCCU (Occurrence); PREP (Preparation); USES (Uses) (amino acid sequence; novel exendin agonist formulations and methods of administration thereof as antidiabetic agents and appetite suppressants) ANSWER 17 OF 20 HCAPLUS COPYRIGHT 2003 ACS 1998:112250 HCAPLUS ACCESSION NUMBER: DOCUMENT NUMBER: 128:192936 Preparation of exendin peptide analogs as TITLE: agonists for regulating gastrointestinal motility Young, Andrew A.; Gedulin, Bronislava; Beeley, INVENTOR(S): Nigel Robert Arnold; Prickett, Kathryn S. Amylin Pharmaceuticals, Inc., USA; Young, Andrew PATENT ASSIGNEE(S): A.; Gedulin, Bronislava; Beeley, Nigel Robert Arnold; Prickett, Kathryn S. SOURCE: PCT Int. Appl., 70 pp. CODEN: PIXXD2 DOCUMENT TYPE: Patent LANGUAGE: English FAMILY ACC. NUM. COUNT: PATENT INFORMATION: KIND DATE APPLICATION NO. DATE PATENT NO. ----**--**\_\_\_\_\_ A1 19980212 WO 1997-US14199 19970808 W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, RW: GH, KE, LS, MW, SD, SZ, UG, ZW, AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG A1 19980225 AU 1997-40636 19970808 AU 9740636 EP 1997-938261 19970808 EP 966297 A1 19991229 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, FI JP 2001501593 Т2 20010206 JP 1998-508263 19970808 US 1996-694954 A 19960808 PRIORITY APPLN. INFO.: WO 1997-US14199 W 19970808 MARPAT 128:192936 OTHER SOURCE(S): Methods for reducing gastric motility and delaying gastric emptying for therapeutic and diagnostic purposes are disclosed which comprise administration of an effective amt. of an exendin or an exendin agonist H-Xaa1-Xaa2-Xaa3-Gly-Thr-Xaa4-Xaa5-Xaa6-Xaa7-Xaa8-Ser-Lys-Gln-Xaa9-Glu-Glu-Glu-Ala-Val-Arg-Leu-Xaa10-Xaa11-Xaa12-Xaa13-Leu-Lys-Asn-Gly-Gly-Xaa14-Ser-Ser-Gly-Ala-Xaa15-Xaa16-Xaa17-Xaa18-Z [Xaa1 = His, Arg, Tyr; Xaa2 = Ser, Gly, Ala, Thr; Xaa3, Xaa7, Xaa12 =

Searcher: Shears 308-4994

independently Asp, Glu; Xaa4, Xaa10 = independently Phe, Tyr,

naphthylalanine; Xaa5, Xaa6 = independently Thr, Ser; Xaa8, Xaa9 = independently Leu, Ile, Val, pentylglycine, Met; Xaa11 = any group Xaa8, tert-butylglycine; Xaa13 = any group Xaa4, Trp; Xaa14-Xaa17 = independently Pro, homoproline, 3-Hyp, 4-Hyp, thioproline, N-alkylglycine, N-alkylpentylglycine, N-alkylalanine; Xaa18 = Ser, Thr, Tyr; Z = OH, NH2; with the proviso that the compd. does not have the formula of exendin-3 or exendin-4] or a pharmaceutically acceptable salt thereof. Methods for treating conditions assocd. with elevated, inappropriate, or undesired post-prandial blood glucose levels are disclosed which comprise administration of an effective amt. of an exendin or an exendin agonist alone or in conjunction with other anti-gastric emptying agents. Thus, exendin-4 acid and [Leu14, Phe25] -exendin-4, prepd. by std. solid-phase methods on a 4-(2,4-dimethoxyphenyl)-Fmocaminomethylphenoxyacetamide norleucine-MBHA resin using 9-fluorenylmethoxycarbonyl (Fmoc)-protected amino acids, inhibited gastric emptying in male HSD rats with EC50 = 0.12 and 0.29 .mu.g. Exendin-4 showed EC50 = 0.27 .mu.g under the same conditions. 130357-25-4P, Exendin-3 (Heloderma horridum) RL: BAC (Biological activity or effector, except adverse); BSU

(Biological study, unclassified); SPN (Synthetic preparation); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES

(prepn. of exendin peptide analogs as agonists for regulating gastrointestinal motility)

THERE ARE 12 CITED REFERENCES AVAILABLE 12 REFERENCE COUNT:

FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 18 OF 20 HCAPLUS COPYRIGHT 2003 ACS 1995:675100 HCAPLUS ACCESSION NUMBER:

123:74913 DOCUMENT NUMBER:

Exendin-3 and exendin-4 polypeptides, and TITLE: pharmaceutical compositions comprising them

Eng, John INVENTOR(S):

PATENT ASSIGNEE(S): USA

U.S., 17 pp. SOURCE: CODEN: USXXAM

DOCUMENT TYPE: Patent English LANGUAGE:

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

APPLICATION NO. DATE PATENT NO. KIND DATE \_\_\_\_ US 1993-66480 19930524 19950613 US 5424286 A 19930524 US 1993-66480 PRIORITY APPLN. INFO.:

This invention encompasses pharmaceutical compns. contg. exendin-3 or exendin-4, fragments thereof, or any combination thereof, and methods for the treatment of diabetes mellitus and the prevention of hyperglycemia.

130357-25-4, Exendin 3 (Heloderma horridum)

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(exendin-3 and exendin-4 polypeptides, and pharmaceutical compns. comprising them)

> 308-4994 Shears Searcher :

ANSWER 19 OF 20 HCAPLUS COPYRIGHT 2003 ACS

ACCESSION NUMBER: 1994:622490 HCAPLUS

DOCUMENT NUMBER: 121:222490

Use of 125I-[Y39]exendin-4 to characterize TITLE:

exendin receptors on dispersed pancreatic acini

and gastric chief cells from guinea pig

AUTHOR(S):

Singh, Gurcharn; Eng, John; Raufman, Jean-Pierre Gastrointestinal Cell Biology Laboratory, State CORPORATE SOURCE:

University of New York-Health Science Center at

Brooklyn, 450 Clarkson Avenue-Box 1196, Brooklyn, NY, 11203-2098, USA

SOURCE: Regulatory Peptides (1994), 53(1), 47-59

CODEN: REPPDY; ISSN: 0167-0115

DOCUMENT TYPE: Journal LANGUAGE: English

We synthesized and iodinated an exendin-4 analog, [Y39]exendin-4 (700 Ci/mmol), for use as a radioligand to characterize exendin receptors on dispersed pancreatic acini and gastric chief cells from guinea pig. Binding of this bioactive radioligand was rapid, temp.-dependent and specific (not inhibited by other pancreatic or gastric secretagogues). Measurement of the ability of exendin-4 to inhibit the binding of 125I-[Y39]exendin-4 indicated the presence of two classes of receptors. Pancreatic acini had 12.5 .times. 1010 binding sites/mg acinar protein of which 6% were high affinity (Kd = 0.5 nM) and 94% were low affinity (Kd = 0.1 .mu.M). Chief cells had 3370 binding sites/cell of which 9% were high affinity (Kd = 0.3 nM) and 91% were low affinity (Kd = 0.2 .mu.M). Washing with 0.2 M acetic acid (pH 2.5), 0.2 M glycine (pH 10.5), or trypsin (100 .mu.g/mL) after 30 min incubation at 37.degree., indicated that 63 and 49% of radioligand was internalized in acini and chief cells, resp. Truncated glucagon-like peptide-1 (tGLP-1), a mammalian peptide sharing 53% homol. with exendin-4, inhibited radioligand binding at the same concns. that altered secretion from acini and chief cells. Glucagon, GLP-1 and GLP-2 inhibited 125I-[Y39] exendin-4 binding only at concns. .gtoreq.100 nM. Exendin(9-39)NH2, a specific exendin-receptor antagonist, potently inhibited 125I-[Y39] exendin-4 binding (IC50 = 6.1 and 3.5 nM in acini and chief cells, resp.). In pancreatic acini and gastric chief cells from guinea pig, exendin-3, exendin-4 and tGLP-1 increase cellular cAMP and modulate enzyme secretion by interacting with high-affinity exendin receptors. 125I-[Y39] exendin-4 is a useful radioligand for studying exendin receptors.

130357-25-4, Exendin 3 (Heloderma horridum) RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); BIOL (Biological study) (cAMP formation and enzyme secretion by pancreas acinus and stomach chief cells response to)

ANSWER 20 OF 20 HCAPLUS COPYRIGHT 2003 ACS ACCESSION NUMBER: 1990:608593 HCAPLUS

DOCUMENT NUMBER: 113:208593

TITLE: Purification and structure of exendin-3, a new

pancreatic secretagogue isolated from Heloderma

horridum venom

AUTHOR(S): Eng, John; Andrews, P. C.; Kleinman, Wayne A.;

Singh, Latika; Raufman, Jean Pierre

CORPORATE SOURCE: Solomon A. Berson Res. Lab., Veterans Aff. Med.

Cent., Bronx, NY, 10468, USA

Searcher : Shears 308-4994

SOURCE: Journal of Biological Chemistry (1990), 265(33),

20259-62

CODEN: JBCHA3; ISSN: 0021-9258

DOCUMENT TYPE: Journal LANGUAGE: English

An assay for Hisl peptides performed by amino-terminal amino acid sequencing was used to screen venom from the Gila monster lizard, H. horridum. Two Hisl peptides were identified: helospectin and a new Hisl peptide that has been named exendin-3 to indicate that it is the third peptide to be found in an exocrine secretion of Heloderma lizards which has endocrine activity, the first two being helospectin (exendin-1) and helodermin (exendin-2). In the lot of H. horridum venom tested, exendin-3 was 5-10-fold more abundant in molar concn. than helospectin. The structure of exendin-3 was analyzed by amino acid sequencing and mass spectrometry. Exendin-3 is a 39-amino acid peptide with a mass of 4200. It contains a carboxyl-terminal amide and has a strong homol. with secretin at its amino-terminal 12 amino acids. The complete structure of exendin-3 is: His-Ser-Asp-Gly-Thr-Phe-Thr-Ser-Asp-Leu-Ser-Lys-Gln-Met-Glu-Glu-Glu-Ala-Val-Arg-Leu-Phe-Ile-Glu-Trp-Leu-Lys-Asn-Gly-Gly-Pro-Ser-Ser-Gly-Ala-Pro-Pro-Pro-Ser. It is 32 and 26% homologous with helospectin and helodermin, resp. It has greatest homol. with glucagon (48%) and human glucagon-like peptide-1 (50%). Exendin-3 (3 .mu.M) stimulated increases in cellular cAMP and amylase release from dispersed guinea pig pancreatic acini.

IT 130357-25-4, Exendin 3 (Heloderma horridum)

RL: PRP (Properties)

(amino acid sequence of)

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Searcher: Shears 308-4994

GenCore version 5.1.4_p5_4578 c) 1993 - 2003 Compugen Ltd.	
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March 28, 2003, 08:25:12; Search time 74 Seconds (without alignments) 70.227 Million cell updates/sec OM protein - protein search, using sw model Run on A

US-09-756-690A-1 208 1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSŠGAPPPS 39 Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

908470 Total number of hits satisfying chosen parameters:

908470 seqs, 133250620 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match, 0% Maximum Match,100% Listing first 45 summaries

A\_Geneseq\_101002:\* Database

/ SIDS2/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:\*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Result		Onerv				
No.	Score	Match	Match Length DB	DB	, QI	Description
-	208	100.0	39	16	AAR80545	Heloderma horridum
7	208	100.0	39	19	AAW61769	Exendin-3, for use
m	208	100.0	39	19	AAW47608	Gila monster exend
4	208	100.0	39	20	AAY31501	Exendin-3 peptide
'n	208	100.0	39	50	AAY03717	Amino acid sequenc
9	208	100.0	39	21	AAB11281	H. horridum exendi
7	208	100.0	39	21	AAB52871	Mexican beaded liz
œ	208	100.0	39	21	AAY94010	Amino acid sequenc
σ	208	100.0	39	22	AAU07378	Glucagon-like pept
10	208	100.0	39	22	AAE08345	Heloderma horridum

90 Pancreatic hormone	31 Mexic	Exendin-	•	32	Gila mons		Gila monster 6	Gila monster				88 exendin agonist pe		Exendin ago	Heloderma suspec	Exendin-4,	09 Gila monster exend		Amino acid	H. suspe	exendin	exendin agonist	exendin agonist	exendin agonist	Extendin-4	Extendin-4	Extendin-4	Extendin-4	Extendin-4 pe	Gila monste	1 Amino acid sequ		0 11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
AAB911	AAB641			-	-	ABB801		ABB07149							AAR80546					-		_	-		AAB52840			AAB52857			AAY94	AAY78	AAU07
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100.0	30	100.0	8	100.0	100.0	00	00	00.	00	00.	9	98.1	98.1	98.1	96.2		96.2		96.2	96.2	96.2	_	96.5	_					96.2	96.2		96.2	
208	208	208	208	208	208	208	208	208	208	208	208	204	204	204	200	200	200	200	200	200	200	200	200	200	200	200	200	200	200	200	200	200	200
11	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	. 34	35	36	37	38	39	40	41	42	43	44	45

# ALIGNMENTS

RES	RESULT
AAR	30545
a	AAR80545 standard; pept1de; 39 AA.
X	
AC	AAR80545;
XX	
Ϊ́	27-FEB-1996 (first entry)
XX	
DE	Heloderma horridum exendin-3.
×	
Κ¥	Exendin-3; diabetes mellitus; hyperglycaemia; insulinotropic pepti
XX	
SO	Heloderma horridum.
×	
Nd	US5424286-A.
×	
δ	13-JUN-1995.
×	
PF	24-MAY-1993; 93US-0066480.
×	
PR	24-MAY-1993; 93US-0066480.
XX	
PA	(ENGJ/) ENG J.
×	
ΡΙ	Eng J;
×	
DR	WPI; 1995-262627/34.
×	
ΡŢ	Stimulating/inhibiting insulin release with exendin polypeptide(s)
PŢ	for treating diabetes mellitus and preventing hyperglycaemia.
XX	
PS >	Claim 5; Columns 13-14; 17pp; English.
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AAR80545 is Heloderma horridum exendin-3. It is an insulinotropic peptide, and can therefore be used in the treatment of diabetes mellitus (types I or II), and for the prevention of hyperglycaemia. It normalises hyperglycaemia through glucose-dependent and insulin-(in)dependent mechanisms.
                                                                                                                                                                                                                                                                                                                                                                                                              Exendin; obesity; type II diabetes; eating disorders; cardiac disease; insulin resistance syndrome; elevated plasma glucose level; agonist.
                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                  Exendin-3, for use in treating disorders related to food intake.
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                                                                                                                          Length 39;
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llarity 100.0%; Pred. No. 9.2e-19;
Conservative 0; Mismatches 0;
                                                                                                                       100.0%; Score 208; DB 16;
1larity 100.0%; Pred. No. 9.2e-19;
Conservative 0; Mismatches 0;
                                                                                                                                                                                Claims 16, 24; Page 8; 214pp; English,
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97US-0055404.
97US-0065442.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bhavsar S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-398796/34
                                                                                                                                     Local Similarity
nes 39; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Heloderma horridum
                                                                                           39 AA;
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08-AUG-1997;
14-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Beeley NRA,
                                                                                                                  Query Match
Best Local S1
Matches 39;
                                                                                           Sequence
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                                                                                                                                                                                                                                                             RESULT 2
AAW61769
ID AAW6
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW47549 describes a generic exendin agonist, provided that it does have the formula of either exendin-3 (AAW47608) or exendin-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Exendin agonists, which reduce gastric motility and delay gastric emptying, can be used to treat spasm (where associated with acute diverticulities or disorders of the bilary tract or sphincter of odd), postprandial dumping syndrome and hyperglycaemia of particularly associated with type 2 diabetes), type 1 diabetes, impaired glucose tolerance, toxin ingestion (an exendin agonist is administered to prevent stomach contents passing into the administered to subjects undergoing gastrointestinal diaphostic investigation, particularly radiological or by magnetic resonance
                                                                                                                                                                                                                                                                            spasm; postprandial dumping syndrome; postprandial hyperglycaemia; type 1 diabetes; impaired glucose tolerance; toxin ingestion; obesity; Gila monster venom; exendin-3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ó;
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                                                                                                                                                                                                                                             Exendin agonist; gastric motility; gastric emptying;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 208; DB 19;
100.0%; Pred. No. 9.2e-19;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Prickett KS, Young AA;
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AAW47608 standard; peptide; 39
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                                                                                                                     (first entry)
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Best Local Similarity 100.0
Matches 39; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AMYL-) AMYLIN PHARM INC
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                                                                                                                                                                                    Gila monster exendin-3.
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(AAW47609).
                                                            AAW47608;
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Gaps

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1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39

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Local Similarity nes 39; Conserv

Query Match Best Loc Matches

Length 39;

Exendin; agonist; diabetes; disorder; plasma glucose; gastric; diagnostic; gastro-intestinal; radiological; generic.

Amino acid sequence of exendin-3.

08-JUN-1999

AAY03717;

AAY03717 standard; peptide; 39 AA.

/note= "C-terminal amide"

Modified-site W09907404-A1

Synthetic

98WO-US16387

06-AUG-1998; 08-AUG-1997; (AMYL-) AMYLIN PHARM INC.

Prickett KS;

seeley NRA,

WPI; 1999-180403/15.

Location/Qualiflers

AAY3150

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RESULT 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Exendin; agonist; GLP-1; glucagon-like peptide; toxic hypervolemia; diuresis; renal plasma flow; glomerular filtration rate; pre-eclampsia; eclampsia of pregnancy; cardiac contractility; renal failure; diuretic; congestive heart failure; nephrotic syndrome; pulmonary edema; cirrhosis; hypertension; urine flow.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   have a low toxicity, and are easily administered intravenously. The present sequence represents an exendin-3 peptide which can be used in the methods of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Increasing urine flow by administering peptides or peptide agonists
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Best Local Similarity 100.0%; Pred. No. 9.2e-19;
Matches 39; Conservative 0; Mismatches 0;
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39
/note= "Citerminal amide"
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                                                                                                                                                                     AAY31501 standard; peptide; 39 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Exendin-3 peptide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
Heloderma horridum.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-AUG-1999
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                                                                                                                                                                                                                                                                        AAY31501;
                                                                       RESULT 4
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xaal.xaa4.xaa6.xaa6.xaa7.xaa8.Ser-Ly8-Gln-Xaa9-Glu-Glu-- Glu-Ala-
yaa3.cly-Thr-Xaa4.xaa5.xaa6.xaa7.xaa13.Leu-Ly8-Asn-Gly-Gly Xaal4.Ser-Ser-
yal-Arg-Leu xaal0. xaal1. xaal2. xaa13.Leu-Ly8-Asn-Gly-Gly Xaal4.Ser-Ser-
yal-Arg-Leu xaal0. xaal1. xaal8.z, whereln xaal 1s His, Arg or Tyr:
Cay Ala. Xaal5. xaa7. xaa8.z, whereln xaa9 is He, Tyr, or
napthylalanine; Xaa5 is Thr or Ser; Xaa6 is Ser or Thr; Xaa7 is Asp or
Clui, Xaa8 is Leu, Ile, Val, pentylglycine, or Met; Xaa9 is Leu, Ile,
pentylglycine, Val, or Met; Xaa10 is Phe, Tyr, or napthylalanine; Xaa15.
Cr Asp; Xaal1 is Trp, Phe, Tyr, or napthylalanine; Xaa15,
and Xaa17 are independently Pro, homoproline, 3Hyp, 4Hyp, thioproline,
n-alkylglycine, N-alkylpentylglycine, or N-alkylalanine; Xaa16,
and Xaa17 are independently Pro, homoproline, 7Hyp, 4Hyp, thioproline,
n-alkylglycine, N-alkylpentylglycine, or N-alkylalanine; Xaa18 is Ser,
nh, or Tyr; and Z is -OH or -NHZ with the proviso that the sequence is
not the amino acid sequences shown in the proviso that above formula where
the specification claims for a second peptide of the above formula where
xaal1 is His, arg, Tyr or 4-imidazoproplonyl. The exendin agonists are
used to treat Type I and II diabetes, disorders which would be benefited by agents useful in delaying and/or slowing gastric emptying.
benefited by agents useful in delaying and/or slowing gastric emptying.
Cradiological examinations, The present sequence represents the amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention relates to exendin agonists which slow gastric emptying lower plasma glucose levels. The peptides are of the formula
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    useful in the treatment of Type I and II

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 208; DB 20;
Pred. No. 9.2e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Delayed gastric emptying is a useful radiological examinations. The prese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Fig 2; 70pp; English.
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Best Local Similarity 100.0%; Pi
Matches 39; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New exendin agonists
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     39 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diabetes
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Length 39; Indels

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The present invention relates to extendins and their agonists which have been modified with molecular weight increasing agents such as polyethylene glycol (PEG). These can be used in the treatment of diabetes, obesity, impaired glucose tolerance, postprandial dumping syndrome, postprandial hyperglycaemia, eating disorders, insulin resistance syndrome, dyslipidaemia and to suppress glucagon secretion.
                                                                                                                                                                                                                              Modified exendin or an exendin agonist linked to one or more polyethylene glycol (PEG) polymers, modulate plasma glucose levels, useful for treating disorders such as diabetes and obesity -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Extendin-3; Gila monster lizard; Mexican Beaded lizard; exen glucagon-like peptide; plasma glucagon; necrolytic erythema; glucagonoma; hyperglucagonemia; diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 208; DB 21; ilarity 100.0%; Pred. No. 9.2e-19; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY94010 standard; pept1de; 39 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amino acid sequence of extendin-3.
                                                                                                                                                                                                                                                                                                            Example 1; Fig 1; 119pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-JAN-1999; 99US-0116380.
30-APR-1999; 99US-0132017.
10-JAN-2000; 2000US-0175365.
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                                                 28-APR-2000; 2000WO-US11814.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (AMYL-) AMYLIN PHARM INC
                                                                                                                      (AMYL-) AMYLIN PHARM INC
                                                                                                                                                             Young A, Prickett K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Young A, Gedulin B;
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                                                                                                                                                                                                WPI; 2000-672834/65.
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Best Local Similarity
Matches 39; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Heloderma horridum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39 AA;
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Modified-site
                                                                                      30-APR-1999;
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           09-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY94010;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel formulation (I) comprising an exendin or exendin agonist peptide, a buffer and an iso-osmolality modifier which has a pl of 3-7. The products of the invention have antidiabetic activity. The exendin or exendin agonist is used to increase the sensitivity of a subject to insulin to treat diabetes and disorders which would benefit from agents which lower plasma glucose levels and disorders which would benefit from agents that delay and/or slow gastric emptying or reducing food intake.
                                                                                                                                                                                                                  Exendin; agonist; treatment; antidiabetic; insulin sensitivity; diabetes; plasma glucose; gastric emptying; food intake.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New formulations comprising an exendin or exendin agonist peptide used for increasing the sensitivity of a subject to insulin to treat diabetes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Extendin; agonist; diabetes; obesity; eating disorder; dyslipidaemia; insulin-resistance syndrome; food intake
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 208; DB 21;
100.0%; Pred. No. 9.2e-19;
iive 0; Mismatches 0;
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                                                                                                                                                                                   H. Morridum exendin 3 peptide SEQ ID NO 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mexican beaded lizard extendin-3 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kolterman O;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Figure 1; 281pp; English.
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                                                                        AAB11281 standard; Peptide; 39 AA.
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                                                                                                                                                                                                                                                                                                                                                                                       10-JAN-2000; 2000US-0116380.
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Best Local Similarity 100.0
Matches 39; Conservative
                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AMYL-) AMYLIN PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Young A, L'Italien JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-514584/46.
                                                                                                                                                                                                                                                                           Heloderma horridum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  39 AA;
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                                                                                                                                                                                                                                                                                                                WO200041546-A2.
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                                                                                                                                       20-FEB-2001
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Length 39;

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Inhibiting antro-duodenal motility, useful to prevent or treat gastrointestinal disorders such as irritable bowel syndrome and non-infectious diarrhoea, comprises administering glucagon-like peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method of inhibiting antro-duodenal motility in a patient, comprising administering a glucagon-like peptide (GLP-1) molecule. The method is used to premedicate or in endoscopic
    Lowering plasma glucagon using exendin, an exendin agonist, a modified exendin or a modified exendin agonist, useful for treating hyperglucagonemia and diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 13; 43pp; English.
                                                                                                                                                                                                                                                                                                                                AAU07378 standard; peptide; 39 AA
                                                Example 1; Fig 1; 96pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-MAR-2000; 2000US-189091P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-MAR-2001; 2001WO-EP02882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Goeke B, Schirra J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-596887/67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GOEK/) GOEKE B.
(SCHI/) SCHIRRA J.
                                                                                                                                                                                                 39 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200168112-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                    18-DEC-2001
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                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                   AAU07378;
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procedures or to treat or prevent non-infectious acute and chronic diarrhoea, post-operative dumping syndrome, irritable bowel syndrome or symptoms associated with narcotics withdrawal. Unlike prior art treatment with glucagon, the invention is not contraindicated in persons with off abetes, does not incur the risks of side effects such as nausea not expensive. The present sequence represents mammalian glucagon-like peptide-1, (GLP-1) homologue, exendin 3 as described in the method of the
                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.0%; Pred. No. 9.2e-19;
Matches 39; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 HSDGTFTSDLSKOMEEEAVRLFIEWLKNGGPSSGAPPPS 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-514422/56.
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                                                                                                                                                                                                                                                                                                39 AA;
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Modified-site
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                                                                                                                                                                                                                                      invention.
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                                                                                                                                                                                                                                The present sequence represents an extendin-3 peptide. Extendins are found in the salivary glands of the Gila monster and Mexican Beaded lizard, and have sequence similarity to glucagon-like peptides. It is useful in the method of the invention. The specification describes a method for lowering plasma glucagon, comprising administering an exendin as one as modified exendin a second, an exendin as modified exendin or a modified exendin useful for lowering plasma glucagon in subjects, preferably humans, suffering from necrolytic erythema or glucagonoma. The method is useful for treating plasma glucagon in subjects, preferably humans, suffering from necrolytic erythema or glucagonoma. The method is also useful for treating hyperglucagonemia and other conditions that would benefit from reduced glucagon levels and/or suppression of glucagon, e.g. type 1 and type 2 dlabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antidiarrhoeic; antlinflammatory; antiaddictive; premedication; antro-duodenal motility; glucagon-like peptide-1; GLP-1; endoscopy; diarrhoea; post-operative dumping syndrome; irritable bowel syndrome; narootics withdrawal; exendin 3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.0%; Pred. No. 9.2e-19;
Matches 39; Conservative 0; Mismatches 0;
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                                                                                                                                                                                            Exendin-3; antilipemic; cardiant; triglyceride; inotropic; dluretic;
coronary heart disease; dyslipidaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The patent discloses a method for modulating plasma or postprandial triglyceride and other lipid levels by administering exendin or an exendin agonist. Exendins have inotropic and diuretic effects. They suppress the secretion of glucagon. Exendin and its agonists have a significant effect on the reduction of blood serum triglyceride concentrations. They are used to treat coronary heart disease and dyslipidaemia, and for modifying postprandial triglyceride levels. The present sequence is exendin-3 peptide from Heloderma horridum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Use of exendin and exendin agonist compounds for modulating triglyceride levels, and treating heart disease and dyslipidemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Score 208; DB 22;
100.0%; Pred. No. 9.2e-19;
ive 0; Mismatches 0;
1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS
                                                                                                                                                                                                                                                                                                 /note= "C-terminal amide"
                                                                                                                                                                       Heloderma horridum exendin-3 peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 10; 161pp; English.
                                                                                                                                                                                                                                                                Location/Qualifiers
39
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                                                                                  AAE08345 standard; peptide; 39
                                                                                                                                                                                                                                                                                                                                                                                                                 10-JAN-2000; 2000US-0175365.
                                                                                                                                                                                                                                                                                                                                                                                    09-JAN-2001; 2001WO-US00719
                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.
nes 39; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                            (AMYL-) AMYLIN PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AA;
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Matches
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Exendin 3; Gila monster venom; GLP-1 analogue; glucagon-like peptide-1; type II diabetees; non-insulin dependent diabetes mellitus; NIDDM; beta-cell function; secretory capacity; impaired glucose tolerance; IGT; beta-cell stimulatory test; diagnostic test; insulinotropic.
                                                                                                                                           Gila monster venom GLP-1 analogue, exendin 3.
                                AAB60252 standard; peptide; 39
                                                                                                                                                                                                                                                                                                                                                                                 14-JUN-2000; 2000WO-US16428
                                                                                                         28-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                              Heloderma suspectum.
                                                                                                                                                                                                                                                                                                                WO200077039-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                   21-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence
                                                                        AAB60252;
RESULT 12
AAB60252
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a modified therapeutic peptide (I) comprising a therapeutically active amino acid region (III) and a reactive group (II) (e.g. succininidy) and matelminod groups) attached to a less therapeutically active amino acid region (IV), which covalently bonds with amino/hydroxyl/thiol groups on blood components to form a peptidase stabilised therapeutic peptide composed of 3-50 amino acids. (I) are useful for modifying therapeutic peptides e.g. hormones, growth factors and neurotransmitters, to protect them from peptidase activity in vivo for the treatment of various disorders. Endogenous therapeutic peptides are not suitable as drug candidates as they require frequent administration due to rapid degradation by peptidases in the body. Modifying and attaching therapeutic peptidases to albumin prevents or reduces the action of peptidases to increase length of activity (half ife) and specificity as bonding to large molecules decreases in the action of the present invention.

Absoluss to Absolute the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modifying and attaching therapeutic peptides to albumin prevents peptidase degradation, useful for increasing length of in vivo activity
                                                                                                                                                                                                                                                                            Protection, endogenous therapeutic peptide; peptidase; conjugation; blood component; modification; succinimidyl; maleimido group; amino; hydroxyl; thiol; hormone; growth factor; neurotransmitter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thibaudeau K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     / Match 100.0%; Score 208; DB 22; Local Similarity 100.0%; Pred. No. 9.2e-19; tes 39; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                            Pancreatic hormone glucagon peptide SEQ ID NO:364
  Holmes DL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 315; 733pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ezrin AM, Milner PG,
                                                                                                                                   AABB1190 standard; Peptide; 39 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0134406.
99US-0153406.
99US-0159783.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-MAY-2000; 2000WO-US13576
                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CONJ-) CONJUCHEM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-112059/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      39 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                 WO200069900-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-MAY-1999;
10-SEP-1999;
15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-NOV-2000.
                                                                                                                                                                                                           22-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                                                                                                                                                     AAB91190;
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Matches
                                                                                                  RESULT 11
                                                                                                                   AAB91190
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The invention relates to a new method for evaluating beta-cell secretory capacity in an individual, or responsiveness of a beta-cell to glucose, comparising the administration of glucose and glucagon-like peptide-1 (GLD-1) or its biologically active analogues. The response in the individual is then measured against the standard response in the individual to determine if the individual has impaired beta-cell function. The method is mestil for detecting impaired beta-cell function. The method is particularly useful for diagnosing impaired of glucose tolerance (IGT) and non-insulin-dependent (type II) diabetes. The method is a rapid test of beta-cell function, which is a marker for impaired glucose tolerance. Onlike prior methods, the method is reliable and without significant adverse side effects and/or patient pain and discomfort. The method also provides information about insulin secretory capacity, and is easy and reproducible. The present sequence represents a gila monster venom GLP-1 analogue peptide referred to in the disclosure
                                                                                                                                                                                                                                                            Evaluating beta-cell secretory capacity and responsiveness to glucose, useful for diagnosing impaired glucose tolerance and diabetes, comprises employing glucagon-like-peptide-1 as a diagnostic test to determine beta-cell function
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Best Local Similarity 100.0%; Pred. No. 9.2e-19;
Matches 39; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 13; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ā
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99US-0333415.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                               (BION-) BIONEBRASKA INC.
                                                                                                                                  Holst JJ, Vilsboll T;
                                                                                                                                                                                                        WPI; 2001-102518/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          39 AA;
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AAB64181
ID AAB64
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AC AAB64
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The invention relates to the use of an exendin (AAB64181-B64182) or an exendin agonist (AAB64185-B64368) for treating gestational diabetes on exendin agonist (AAB64185-B64368) for treating gregnancy, and is due to a combination of increased insulin resistance and a diminished ability to increase insulin resistance and insulin secretion increase. GDM pregnancies are associated with complications in both the mother and the pregnancies are associated with complications in both the mother and the ingernancies are associated with complications in both the mother and the infections. GDM results in an elevated rates of Caesarian delivery, in a neural tube defects, and is associated with an increased risk of the contrabilities such as neural tube defects, and is associated with an increased risk of the contrabilities such as hypoglycaemia, hypocalcaemia, and subsequent chlidhood and adolescent obesity. Exendins are peptides from the salivary secretions of the Glim monster (exendin-4) and the Mexican beaded lizard (exendin-3) which exhibit homology with several members of the glucagon-like peptide family, particularly GLP-1, and have similar contraindicated for GDM, exendins and exendin contraindicated for GDM, exendin section of the exercise exerce prolonged effect on hypoglycaemial and the measur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Use of exendins or exendin agonists for lowering or reducing blood glucose levels and treating gestational diabetes mellitus in a subject, especially in a human
                                                               Exendin-3; gestational diabetes mellitus; GDM; insulin resistance;
pregnancy complication; neonatal abnormality; blood glucose modulator;
insulinotropic; anorectic; Mexican beaded lizard.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       which is specifically claimed for use in the invention
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100.0%; Score 208; DB 22;
Best Local Similarity 100.0%; Pred. No. 9.2e-19;
Matches 39; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
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                      Mexican beaded lizard exendin-3, SEQ ID NO:1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 11; Page 10-11; 133pp; English.
                                                                                                                                                                                                                                                                                                                                     23-MAY-2000; 2000WO-US14231.
                                                                                                                                                                                                                                                                                                                                                                                   99US-0323867.
                                                                                                                                                                                                                                                                                                                                                                                                                                     (AMYL-) AMYLIN PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hiles R, Prickett KS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-137634/14.
                                                                                                                                                                                !@loderma horridum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 39 AA;
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                                                                                                                                                                                                                                                                                                                                                                                      01-JUN-1999;
                                                                                                                                                                                                                                                                                07-DEC-2000.
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AAB48800
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The invention relates to modified insulinotropic peptides (ITP9), or derivatives thereof which comprise a reactive group which reacts with anino groups, hydroxyl groups or thich groups on blood components (e.g., serum albumin) to form a stable covalent bond. The insulinotropic serum albumin) to form a stable covalent bond. The insulinotropic or serum albumin and contain a reactive group such as a malehmido (51b-1) or exendin and contain a reactive group such as a malehmido or succinimidyl group. The peptides of the invention act by stimulating the synthesis or expression of insulin. A composition comprising a peptide of the invention is useful for treating diabetes, paticularly type II (maturity onset) diabetes. It is also useful as a saticularly type II (maturity onset) diabetes. It is also useful as a settle paticularly type II (maturity onset) diabetes. It is also useful as a settle paticularly type II (maturity onset) diabetes. It is also useful as a contient the treatment of nervous system disorders including anxiety, psychosis, seizures, panic attacks, hysteria and sleep considers; to induce an anxiolytic effect on the central nervous system (CNS); to activate the CNS for the treatment of disorders such as conjugation of a pertide of the invention to a long surgery. The conjugation of a peptide of the invention to a long surgery. The conjugation of a peptide of the invention therefore the processor of peptidases. The peptides of the invention therefore proteolytic degradation. The present sequence represents an eless susceptible to insulinotropic peptide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                              Insulinotropic peptide; insulin production; GLP-1 derivative; glucagon-like peptide 1; exendin derivative; reactive group; peptidase stabilisation; blood protein; conjugation; type II diabetes; insulin resistance; nervous system disorder; sedative; anxiolytic; antidiabetic; neuroprotective; tranquiliser; anticonvulsant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel modified insulinotropic peptides for treating diabetes, nervous system disorders and for post surgery treatment, has reactive groups which react with amino, hydroxy or thiol groups on blood components
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leblanc A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Holmes DL,
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llarity 100.0%; Pred. No. 9.2e-19;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   | HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ezrin AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; Page 88-89; 96pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    L'Archeveque B,
                                                                                                                                                                                                                                                                                                      17-MAY-2000; 2000WO-US13563.
                                                                                                                                                                                                                                                                                                                                              99US-0134406.
99US-0159783.
               Exendin-3, SEQ ID NO:11.
                                                                                                                                                                                                                                                                                                                                                                                                            (CONJ-) CONJUCHEM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-025008/03.
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Best Local Similarity
Matches 39; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         39 AA;
                                                                                                                                                                                                                          WO200069911-A1.
                                                                                                                                                                                                                                                                                                                                                  17-MAY-1999;
                                                                                                                                                                               Unidentified.
                                                                                                                                                                                                                                                                                                                                                                  15-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             St Pierre S;
                                                                                                                                                                                                                                                             23-NOV-2000.
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AAB36419
ID AAB36.
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AC AAB36
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AAB48800 standard; pept1de; 39

09-MAR-2001

AAB48800;

Location/Qualifiers 39

/note= "amidated"

99US-0303016

us-09-756-690a-1.rag

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The present invention describes a method for the amelioration of brain tissue injury caused by reperfusion of blood flow comprising administering a composition including a compound which binds to a maintistering a composition including a compound which binds to a ceceptor for glucagon-like peptide-l (GLP-l), in a pharmaceutical carrier. The method is used for amelioration of brain tissue injury caused by reperfusion of blood flow following a period of ischaemia. Cup-l is used for metabolic intervention to improve the function of ischaemic and reperfused brain cells. It treats patients after an acute stroke or haemorinage and tissue damage arising from a medical procedure intoke or haemorinage and tissue damage arising from a medical procedure involving a reperfusion event. GLP-l is an ideal alternative to insulin for the treatment of stroke as it stimulates endogenous to insulin for the treatment of stroke as it stimulates endogenous to insulin for the treatment of normo- to hypersyloaemia but not during hypoglycaemia, thus protecting against the development of severe brain anabolism, enhancing insulin effectiveness by suppressing glucagon and maintains euglycaemia or mild hypoglycaemia within no risk of severe hypoglycaemia. The present sequence represents a Glia monster venom ceptide which is homologous to GLP-l, and is glven in the
                                                                       Glucagon-like peptide-1; GLP-1; GLP-2; vasotropic; cerebroprotective; brain tissue injury; reperfusion; blood flow; ischaemia; acute stroke; metabolic intervention; haemorrhage; tissue damage; medical procedure; surgical; insulin; hyperglycaemia; hypoglycaemia; brain anabolism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A method for amelioration of brain tissue injury comprises administering a composition including a compound which binds to receptor for glucagon-like peptide-1
                                     Gila monster venom exendin 3 peptide SEQ ID NO:7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 8; 19pp; English
                                                                                                                                                                                                                                                                                                                                                                                                         01-MAY-2000; 2000WO-US11652.
28-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Coolidge TR, Ehlers MRW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BION-) BIONEBRASKA INC.
                                                                                                                                                                                                          Heloderma suspectum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-015911/02.
                                                                                                                                                                                                                                                                                                                                WO200066142-A2
                                                                                                                                                                                                                                                                 Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-APR-1999;
                                                                                                                                                                euglycaemia.
                                                                                                                                                                                                                                                                                                                                                                     09-NOV-2000
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Gaps ö Length 39; Indels 100.0%; Score 20%; DB 22; 100.0%; Pred. No. 9.2e-19; rative 0; Mismatches 0; Query Match Best Local Similarity 100. Matches 39; Conservative

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Search completed: March 28, 2003, 08:37:20 Job time: 75 secs

42, Appli 1, Appli 4, Appli 8, Appli 1, Appli 13, Appli

Run on 14

Sequence:

Database

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Sequence 1. Application US/08066480
Patent No. 542426
GENERAL INFORMATION:
APPLICANT: Eng, John
TITLE OF INVENTION: Pharmaceutical Compositions And Use of
TITLE OF INVENTION: Exendin-3 and Exendin-4 for Treatment of Diabetes Mellitus
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                              Sequence 1
Sequence 5
Sequence 6
Sequence 6
Sequence 6
Sequence 7
Sequence 1
Sequence 2
Sequence 3
Sequence 3
                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESS.
STREET: 10 S. Wacker. --
STREET: 10 S. Wacker. --
STREET: 11110018
COUNTRY: 11110018
COUNTRY: 126606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM FC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: 08/08/066,480
FILING DATE: 24 HAR-1993
FLING DATE: 24 HAR-1993
CLASSIFICATION: 14 HORMATION:
NAME: MCCONDIGHT, John J
REGISTRATION NUMBER: 26,949
REFERENCE/DOCKET NUMBER: 29,084
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION 11:
TELEPHONE: 312-715-1100
TELECOMMUTCATION 100: 1:
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100.0%; Score 20%; DB 1;
Best Local Similarity 100.0%; Pred. No. 7.1e-20;
Matches 39; Conservative 0; Mismatches 0;
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US-09-623-618B-13

US-09-623-618B-20

US-09-623-618B-21

US-08-623-618B-21

US-08-623-618B-21

US-08-623-618B-21

US-08-623-618B-21

US-09-209-799D-14

US-09-208-750-1

US-09-258-750-4

US-09-258-750-4

US-09-258-750-8

US-09-288-750-1

US-09-288-750-1

US-09-288-711-1

US-09-289-111-1

US-09-398-111-1

US-09-398-111-1

US-09-398-111-1

US-09-398-111-4

US-09-398-111-4

US-09-398-111-4

US-09-398-111-4

US-09-398-111-4

US-09-398-111-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Allegretti & Witcoff, Ltd. STREET: 10 S. Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 312-715-1234
INFORMATION FOR SEX ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 39 amino acids
TYPE: amino acid
STRANDEDNESS: single
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     NAME/KEY: Peptide
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Sequence 9, Al
Sequence 12,
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Sequence 18,
Sequence 31,
Sequence 32,
Sequence 5, A
Sequence 6, A
Sequence 8, A
Sequence 8, A
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Sequence 1
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Sequence 3
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                                                                                                                                                                                                                                                                                                                       US-09-756-690A-1
208
1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
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'cgn2_b/ptdata1/liaa/5A_COMB.pep:*

'cgn2_b/ptdata1/liaa/5B_COMB.pep:*

'cgn2_b/ptdata1/liaa/6A_COMB.pep:*

'cgn2_b/ptdata1/liaa/Ba_COMB.pep:*

'cgn2_b/ptdata1/liaa/Pa_COMB.pep:*

'cgn2_b/ptdata1/liaa/Pa_COMB.pep:*

'cgn2_b/ptdata1/liaa/Pa_COMB.pep:*
                                      GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-302-596-7
US-09-302-596-7
US-09-303-415-7
US-09-303-016-7
US-09-623-618B-19
US-09-623-618B-34
US-09-623-618B-34
US-09-303-618-34
US-09-303-618-18
US-09-303-618-18
US-09-623-618B-12
US-09-623-618B-12
US-09-623-618B-18
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US-09-303-415-8
US-09-303-016-8
US-09-623-618B-14
US-09-303-016-8
US-09-623-618B-14
US-09-623-618B-14
US-09-623-618B-14
US-09-623-618B-14
US-09-623-618B-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                262574 seqs, 29422922 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                     protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 2000000000
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Gaps

1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39

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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Coolidge, Thomas R.
APPLICANT: Coolidge, Thomas R.
APPLICANT: Coolidge, Thomas R.
APPLICANT: Coolidge, Matio R.W.
TITLE OF INVENTION: Metabolic Intervention with GLP-1 or its Biologically
TITLE OF INVENTION: Active Analogues to Improve the Function of the
TITLE OF INVENTION: Active Analogues to Improve the Function of the
TITLE OF INVENTION: 1sohemic and Reperfused Brain
FILE REPERBACE: 1904-043.30
CURRENT FILING DATE: 1999-04-30
FRIOR APPLICATION NUMBER: 60/103,498
FRIOR FILING DATE: 1998-10-08
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO?
SEQ ID NO?
                                                                                                                                                                                        APPLICANT: Holst, Jens J.
APPLICANT: Holst, Jens J.
APPLICANT: Holst, Tina
TITLE OF INVENTION: GLP-1 as a Diagnostic Test to Determine Beta-Cell
TITLE OF INVENTION: Function and the Presence of the Condition of IGT and
TITLE OF INVENTION: Type II Diabetes
FILE REPERENCE: P039870S0
CURRENT APPLICATION NUMBER: US/09/333,415
CURRENT FILING DATE: 1999-06-15
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 39;
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Best Local Similarity 100.0%; Pred. No. 7.1e-20;
Matches 39; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 208; DB 4;
100.0%; Pred. No. 7.1e-20;
Live 0; Mismatches 0;
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Patent No. 6429197
                                                                                                                     Sequence 7, Application US/09333415 Patent No. 6344180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Heloderma suspectum US-09-303-016-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: PRT
; ORGANISM: Heloderma suspectum
US-09-333-415-7
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APPLICANT: L'Archeveque, Ben
APPLICANT: Eziln, Alan M.
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Best Local Similarity 100.
Matches 39; Conservative
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US-09-623-618B-19
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US-09-303-016-7
                                                                                                      US-09-333-415-7
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APPLICANT: Coolidge, Thomas R.
TITLE OF INVENTION: Metabolic Intervention with GLP-1 to Improve the Function of TITLE OF INVENTION: Ischemic and Reperfused Tissue
FILE REFERENCE: P03660013
CURRENT APPLICATION NUMBER: US/09/302,596
CURRENT FILING DATE: 1999-04.30
PRIOR APPLICATION NUMBER: 60/103,498
PRIOR PILING DATE: 1999-10-08
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
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100.0%; Score 208; DB 4; Length 39;
Best Local Similarity 100.0%; Pred. No. 7.1e-20;
Matches 39; Conservative 0; Mismatches 0; Indels
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APPLICANT: L'Archeveque, Benoit
APPLICANT: Ezin, Alan M.
APPLICANT: Leblanc, Anouk
APPLICANT: Long LASTING INSULINOTROPIC PEPTIDES
FILE REFERENCE: 500802001620
CURRENT APPLICATION NUMBER: US/09/623,618B
CURRENT FILING DATE: 2000-09-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 208; DB 4; Best Local Similarity 100.0%; Pred. No. 7.1e-20; Matches 39; Conservative 0; Mismatches 0;
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| HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
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PRIOR APPLICATION NUMBER: PCT/US00/13563
PRIOR PILING DATE: 2000-05-17
PRIOR APPLICATION NUMBER: 60/159,783
PRIOR APPLICATION NUMBER: 60/159,783
PRIOR APPLICATION NUMBER: 60/134,406
PRIOR PILING DATE: 1999-05-17
NUMBER OF SEQ ID NOS: 35
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11
LENGTH: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11, Application US/09623618B
Patent No. 6329336
                                                                                                                                                            Sequence 7, Application US/09302596 Patent WG. 6284725 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Gila Monster venom
US-09-302-596-7
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Sequence 2, Application US/08066480
Sequence 2, Application US/08066480
Patent No. 542426
GENERAL INFORMATION:
APPLICANT: Eng, John
TITLE OF INVENTION: Pharmaceutical Compositions And Use of
TITLE OF INVENTION: Exendin-3 and Exendin-4 for Treatment of Diabetes Mellitu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Peptide NAMEKEY: MOD_RES LOCATION: 33 LOCATION: 33 CHER INFORMATION: Xaa represents Lys(E-AEEA-AEEA-MPA)-NH2-5TFA and where "E" US-09-623-618B-34
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FILE OF INVENTION: LONG LASTING INSULINOTROPIC PEPTIDES
FILE REFERENCE: 50066201620
CURRENT APPLICATION NUMBER: US/09/623,618
CURRENT FILING DATE: 2000-09-05
PRIOR APPLICATION NUMBER: PCT/US00/13563
PRIOR FILING DATE: 2000-05-17
PRIOR PLICATION NUMBER: 60/159,783
PRIOR FILING DATE: 1999-10-15
PRIOR PLING DATE: 1999-10-15
PRIOR PLING DATE: 1999-61-15
NUMBER OF SEQ 1D NOS: 35
SOFTWARE: FastSEQ for Windows Version 4.0
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.0%; Pred. No. 7.4e-20;
Matches 39; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
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     NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: Allegretti & Witcoff, Ltd.
STREET: 10 S. Wacker Drive
CITY: Chicago
STATE: 111inois
COUNTRY: USA
ZIP: 60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UMBER: US/08/066,480
24-MAR-1993
                                                                                                                                                     Sequence 34, Application US/09623618B Patent No. 6329336
                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Bridon, Dominique P.
APPLICANT: L'Archeveque, Benoit
APPLICANT: Ezin, Alan M.
APPLICANT: Holmes, Darren L.
APPLICANT: Leblanc, Anouk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                          RESULT 8
US-09-623-618B-34
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LENGTH: 40
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US-09-623-618B-33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Peptide US-09-623-618B-19
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GENERAL INFORMATION:
APPLICANT: Bridon, Dominique P.
APPLICANT: Ezin, Alan M.
APPLICANT: Ezin, Alan M.
APPLICANT: Ezin, Alan M.
APPLICANT: Holbanc, Anouk
APPLICANT: Holbanc, Anouk
APPLICANT: Holbanc, Anouk
APPLICANT: Leblanc, Anouk
APPLICANT: St. Plarre, Serge
TITLE OF INVENTION: LONG LASTING INSULINOTROPIC PEPTIDES
FILE REFERENCE: 500862001620
CURRENT PAPLICATION NUMBER: US/09/623,618B
CURRENT PRILNG DATE: 2000-09-05
PRIOR APPLICATION NUMBER: EXTUSSO/1363
PRIOR FILING DATE: 1999-10-15
PRIOR FILING DATE: 1999-10-15
PRIOR PRILING DATE: 1999-05-17
NUMBER OF SEQ ID NOS: 35
SEQ ID NOS: 35
SEQ ID NOS: 35
LENGTH: 40
APPLICANT: Holmes, Darren L.
APPLICANT: Leblanc, Anouk
APPLICANT: Leblanc, Anouk
APPLICANT: Leblanc, Anouk
APPLICANT: Serge
TITLE OF INVENTION: LONG LASTING INSULINOTROPIC PEPTIDES
FILE REFERENCE: 500862001620
CURRENT PELICATION NUMBER: US/09/623,618B
CURRENT PILING DATE: 2000-09-05
PRIGR APPLICATION NUMBER: PCT/US00/13563
PRIGR APPLICATION NUMBER: 60/134,406
PRIOR FILING DATE: 1999-10-15
PRIOR FILING DATE: 1999-05-17
NUMBER OF SEQ ID NOS: 35
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 40
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Best Local Similarity 100.0%; Pred. No. 7.4e-20;
Matches 39; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
100.0%; Score 20%; DB 4;
Best Local Similarity 100.0%; Pred. No. 7.4e-20;
Matches 39; Conservative 0; Mismatches 0;
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ORGANISM: Artificial Sequence
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Sequence 9, Application US/09333415
Sequence 9. Application US/09333415
Patent No. 6344180
GENERAL INFORMATION:
APPLICANT: HOIST, Jens J.
APPLICANT: Vilaboll, Tina
TITLE OF INVENTION: GLP-1 as a Diagnostic Test to Determine Beta-Cell
TITLE OF INVENTION: Pupe-II Diabetes
FILE REPERBNCE: P03987US0
CURRENT APPLICATION NUMBER: US/09/333,415
CURRENT FILING DATE: 1999-06-15
NUMBER OF SEQ ID NOS: 13
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GENERAL INFORMATION:
APPLICANT: COolidge, Thomas R.
APPLICANT: Colidge, Thomas R.
TITLE OF INVENTION: Metabolic Intervention with GLP-1 or its Biologically
TITLE OF INVENTION: Active Analogues to Improve the Function of the
TITLE OF INVENTION: Ischemic and Reperfused Brain
FILE REPERRONCE: PO3660082
CURRENT APPLICATION NUMBER: US/09/303,016
PRIOR APPLICATION NUMBER: 60/103,498
PRIOR FILING DATE: 1999-04-30
PRIOR FILING DATE: 1998-10-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Pred. No. 7.4e-19;
1; Mismatches 1; Indels
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96.2%; Score 200; DB 4;
Best Local Similarity 94.9%; Pred. No. 7.4e-19;
Matches 37; Conservative 1; Mismatches 1
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PRIOR APPLICATION NUMBER: PCT/US00/13563
PRIOR FILING DATE: 2000-05-17
PRIOR PLING DATE: 1999-05-17
PRIOR PLING DATE: 1999-10-15
PRIOR APPLICATION NUMBER: 60/134,406
PRIOR PILING DATE: 1999-05-17
NUMBER OF SEQ ID NOS: 35
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12
LENGTH: 39
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US-09-303-016-9
Sequence 9, Application US/09303016
; Patent No. 6429197
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                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 96.2%;
Best Local Similarity 94.9%;
Matches 37; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PatentIn Ver. 2.0
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SEQ ID NO 9
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Patent No. 6284725
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: COLIDGE, Mario R.W.
APPLICANT: Ehlers, Mario R.W.
APPLICANT: Ehlers, Mario R.W.
ITILE OF INVENTION: Metabolic Intervention with GLP-1 to Improve the Function of TILE OF INVENTION: Metabolic and Reperfused Tissue
FILE REFERENCE: P03660051
CURRENT FAPLICATION NUMBER: US/09/302,596
CURRENT FILING DATE: 1999-04-30
PRIOR PLICATION NUMBER: 60/103,498
PRIOR PLICATION NUMBER: 60/103,498
PRIOR FILING DATE: 1998-110-08
NUMBER OF SEQ ID NOS: 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     96.2%; Score 200; DB 1; Length 39; 94.9%; Pred. No. 7.4e-19; 1. Mismatches 1; Indels
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APPLICANT: Leblanc, Anouk
APPLICANT: St. Pleire, Serge
ITILE OF INVENTION: LONG LASTING INSULINOTROPIC PEPTIDES
FILE REFERENCE: 500862001620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 96.2%; Score 200; DB 4; Best Local Similarity 94.9%; Pred. No. 7.4e-19; Matches 37; Conservative 1; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; NAME/KEY: Peptide
; LOCATION: 1..39
; OTHER INFORMATION: /label= Exendin-4
US-08-066-480-2
                                    NAME: McDonnell, John J
REGISTRATION NUMBER: 26,949
REFERENCE/TOKET NUMBER: 93,084
TELECOMMUNICATION INFORMATION:
TELEPONE: 312-715-1000
TELEFENX: 312-715-1234
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 39 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 12, Application US/09623618B Patent No. 6329336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Bridon, Dominique P.
APPLICANT: L'Archeveque, Benoit
APPLICANT: Ezrin, AAlan M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Gila Monster venom
US-09-302-596-9
                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 94.9
Matches 37; Conservative
                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-623-618B-12
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                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
                                                                                                                            0;
                                                                                                                                                                                                                                    Sequence 3.2 Application US/09623618B
Fatent No. 6329336
GENERAL INFORMATION:
APPLICANT: Bridon, Dominique P.
APPLICANT: Eratio, Alan M.
APPLICANT: Holmes, Darren L.
APPLICANT: APPLICANT: St. Plerre, Serge
TITLE OF INVENTION: LONG LASTING INSULINOTROPIC PEPTIDES
FILE REFERENCE: 500862001620
CURRENT APPLICATION NUMBER: US/09/623,618B
CURRENT FILING DATE: 2000-09-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
96.2%; Score 200; DB 4; Length 40;
Best Local Similarity 94.9%; Pred. No. 7.6e-19;
Matches 37; Conservative 1; Mismatches 1; Indels
                                                                                                  Length 39;
                                                                                                                              Indels
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                                                                                                                                                                     96.2%; Score 200; DB 4;
illarity 94.9%; Pred. No. 7.4e-19;
Conservative 1; Mismatches 1
                                                                                                                                                       1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                ) ORGANISM: Heloderma suspectum
US-09-303-016-9
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 9
                                                                                                     Query Match
Best Local Similarity
Matches 37; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 15
US-09-623-618B-31
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US-09-623-618B-18
                                                   TYPE: PRT
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; OTHER INFORMATION: Xaa represents Lys(E-MPA)-NH2-5TFA and where "E" represents US-09-623-618B-31
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                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Peptide
NAME/KEY: MOD_RES
LOCATION: 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 HSDGTFTSDLSKOMEEEAVRLFIEWLKNGGPSSGAPPPS 39
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PRIOR APPLICATION NUMBER: PCT/US00/13563
PRIOR FILING DATE: 2000-05-17
PRIOR APPLICATION NUMBER: 60/159,783
PRIOR FILING DATE: 1999-10-15
PRIOR FILING DATE: 1999-10-15
PRIOR FILING DATE: 1999-05-17
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 31
LENGTH: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Search completed: March 28, 2003, 08:40:49 Job time: 29 secs
                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    96.2%;
94.9%;
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Best Local Similarity 94.9%
Matches 37; Conservative
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March 28, 2003, 08:37:28.; Search time 14 Seconds (without alignments) 163.587 Million cell updates/sec
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2: /cgn2_6/ptodata/1/pubpaa/PUS06_NEW_PUB.pep:*

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14: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
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208
1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSŠGAPPPS 39
          GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                       237916 seqs, 58723674 residues
                                                                                                                                                                                                                                                                                Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                        OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 2000000000
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                                                                                                                           Title:
Perfect score:
                                                                                                                                                  Sequence:
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                                                                              on
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## Sequence 7, Appli Sequence 7, Appli Sequence 1, Appli Sequence 7, Appli Sequence 7, Appli Sequence 19, Appli Sequence 13, Appl Sequence 14, Appl Sequence 14, Appl Sequence 12, Appli Sequence 2, Appli Sequence 2, Appli Sequence 9, Appli Sequence 2, Appli Sequence 1, Appli Description Sequence US-09-876-388-11 US-09-851-738-7 US-09-851-738-7 US-09-851-738-7 US-09-851-738-7 US-09-961-11 US-09-951-0218-7 US-09-951-0218-7 US-09-876-388-19 US-09-876-388-33 US-09-756-690A-14 US-09-756-690A-2 US-09-756-690A-2 US-09-876-388-12 US-09-756-690A-1 SUMMARIES a 100 100 100 100 100 100 100 100 Length % Query Match Score Result No.

96.2         39         10         US-09-982-978-9         Sequence 9, Appli           96.2         40         10         US-09-953-021B-9         Sequence 9, Appli           96.2         40         10         US-09-876-388-18         Sequence 31, Appl           96.2         40         10         US-09-876-388-31         Sequence 32, Appl           95.7         39         US-09-756-690A-12         Sequence 25, Appl           94.7         39         US-09-756-690A-18         Sequence 10, Appl           94.7         39         US-09-756-690A-18         Sequence 10, Appl           94.7         39         US-09-756-690A-18         Sequence 25, Appl           94.7         39         US-09-03-869-10         Sequence 10, Appl           94.7         39         US-09-03-869-10         Sequence 29, Appl           94.7         39         US-09-03-869-18         Sequence 29, Appl           94.7         39         US-09-03-869-18         Sequence 29, Appl           94.7         39         US-09-03-869-18         Sequence 29, Appl           94.2         39         US-09-03-869-13         Sequence 13, Appl           94.2         39         US-09-03-869-13         Sequence 13, Appl           <	ALIGNMENTS  100 US/09756690A  100 018504A1  100 US/09756690A  100 US/09756690A  100 US/09756690A  100 US/09756690A  100 US/09756690A  100 US/097756690A  100 US/097756690A  100 US/09775690A  100 US/09775965  100 US/097596690A  100 US/09759690A  100 US/0975690A  100 US/09756A  100 US/09756A  100 US/09756A  100 US/09756A  100 US/0	tch 100.0%; Score 208; DB 9; Length 39; al Similarity 100.0%; Pred. No. 1.4e-19; 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0; 39; Conservative 0; Mismatches 39 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39 11, Application US/09876388 0. US20020049153a1 INFORMATION: INT: Briddon, Dominique P. NT: L'Archeveque, Benoit.
0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	690A-1 1, Application No. 1906 1, No. No.	h, Similarii Similarii DGTFTSDLSi H          HGTFTSDLSi H8-11 US200200 US200200 CS-11 CS-1
2000 2000 2000 2000 2000 2000 2000 200	ULT 1  09-756-690A-1  equence 1, Applica  ublication No. US2  MAPLICANT: VOLTERM  APPLICANT: KOLTERM  APPLICANT: KOLTERM  APPLICANT: KOLTERM  APPLICANT: KOLTERM  APPLICANT: COUNGY,  TITLE OF INVENTION  FILE REFERENCE 24  CURRENT FILING DAT  FRICH REPLICATION  CURRENT FILING DATE  NUMBER OF SEQ ID N  ELENGTH: 39  LENGTH: 39  LENGTH: 39  TYPE: PRT  CREANISM: Heloder  REALIURE:  OTHER INFORMATION  FRATURE:  OTHER INFORMATION  COTHER INFORMATION  COTHER INFORMATION  OTHER INFORMATION	
04044444444444444444444444444444444444	RESULT Seque Publi General Publi APPL TITL TITL FILE CURR NUMB NUMB SEO I SEO	Query Me Best Loc Matches Oy 1 Db 1 Db 7 RESULT 2 US-09-876 ; Sequence ; GENERAL ; APPLICA

Gaps

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APPLICANT: COGLIDGE, THOMAS R.
APPLICANT: COCLIDGE, THOMAS R.
APPLICANT: EHLERS, MARIO
TITLE OF INVENTION: TREATHENT OF ACUTE CORONARY SYNDROME WITH GLP-1
FILLS REFERENCE: 089187/0395
CURRENT TILLNG DATE: 2001-05-18
PRIOR APPLICATION NUMBER: 60/205,239
PRIOR PILING DATE: 2000-05-19
APPLICANT: EHLERS, MARIO
TITLE OF INVENTION: TREATMENT OF ACUTE CORONARY SYNDROME WITH GLP-1
FILLE REFERENCE: 089187/0395
CURRENT PILING DATE: 2001-03-14
PRIOR RELING DATE: 2001-05-18
PRIOR FILING DATE: 2001-05-18
SOFTWARE: PALENTIN VOS: 13
SOFTWARE: PALENTIN VET: 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 39;
                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Unknown Organism: Exendrin US-09-805-507-7
                                                                                                                                                                                                                                                                                                                                                                                                                      Length 39;
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Pred. No. 1.4e-19;
; Mismatches 0;
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Best Local Similarity 100.0%; Pred. No. 1.4e-19;
Matches 39; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-859-804-7; Sequence 7, Application US/09859804; Sequence No. US-0020107206Al patent No. US-0020107206Al parent INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 39; Conservative 0;
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                                                                                                                                                                                                                                                                                                  ORGANISM: Unknown Organism
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SOFTWARE: Patentin Ver. SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 6
US-09-003-869-1
                                                                                                                                                                                                                                                                                                                                FEATURE:
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Patent No. US20020055460A1

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Coolidge, Thomas R.
APPLICANT: Enlers, Mario R.W.
TITLE OF INVENTION: Metabolic Intervention with GLP-1 to Improve the Function of TITLE OF INVENTION: Ischemic and Reperfused Tissue
FILE REPERENCE: P03660US1
CURRENT APPLICATION NUMBER: US/09/851,738
CURRENT FILING DATE: 201-05-09
PRIOR PILING DATE: 201-05-09
PRIOR PILING DATE: 1999-04-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 39;
                                                     APPLICANT: Lebland, Anouk
APPLICANT: St. Plerre, Serge
ITLE OF INVENTION LONG LEASTING INSULINOTROPIC PEPTIDES
FILE REFERENCE: 500862001610
CURRENT APPLICATION NUMBER: US/09/876,388
CURRENT FILING DATE: 2001-09-24
PRIOR|FILING DATE: 2000-09-05
PRIOR|FILING DATE: 2000-09-05
PRIOR|FILING DATE: 1909-05-17
PRIOR|FILING DATE: 1999-05-17
PRIOR|FILING DATE: 1999-05-17
PRIOR FILING DATE: 1999-05-17
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PRIOR FILING DATE: 1999-05-17
SPRIOR FILING DATE: 1999-05-17
SPRIOR FILING DATE: 1999-05-17
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PRIOR FILING DATE: 1999-05-17
SEC ID NO 11
LENGTHRE: PATCHTON VOWER: 60/134,406
PRIOR FILING DATE: 1999-05-17
LENGTHRE: PATCHTEN VOWER: 2.1
SEC ID NO 11
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Pred. No. 1.4e-19;
); Mismatches 0;
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illarity 100.0%; Pred. No. 1.4e-19;
Conservative 0; Mismatches 0;
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100.0%; Score 20%; Dest Local Similarity 100.0%; Pred. No. 1.4
Matches 39; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: PRT; ORGANISM: Gila Monster venom US-09-851-738-7
             Ezrin, Alan M.
Holmes, Darren L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 7
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Best Local Similarity
Matches 39; Conserv
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Sequence 1, Application US/09003869A

Patent No. US20020137666A1

GENERAL INFORMATION:

APPLICANT: BEELEY, NIGEL ROBERT ARNOLD

APPLICANT: BELEZ, NIGEL ROBERT ARNOLD

TITLE OF INVENTION: USE OF EXENDINS AND AGONISTS THEREOF FOR

TITLE OF INVENTION: UTE REDUCTION OF FOOD INTAKE

FILE REFERENCE: 231/181

CURRENT FILING DATE: 1998-01-07

EARLIER APPLICATION NUMBER: US 60/034,905

EARLIER FILING DATE: 1997-01-07

Gaps

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US-98/0-3188-19
Sequence 19, Application US/09876388
Patent No. US20020049153A1
GENERAL INFORMATION:
APPLICANT: Bridon, Dominique P.
APPLICANT: Ezin, Alan M.
APPLICANT: Ezin, Alan M.
APPLICANT: Lablanc, Anan M.
APPLICANT: OF INVENTION: LONG LASTING INSULINOTROPIC PEPTIDES
TILE REFERENCE: 500862001610
CURRENT APPLICATION NUMBER: 02/09/876,388
CURRENT FILING DATE: 2001-09-24
PRIOR FILING DATE: 2000-09-05
PRIOR FILING DATE: 1999-10-15
PRIOR FILING DATE: 1999-10-15
PRIOR FILING DATE: 1999-10-15
PRIOR FILING DATE: 1999-05-17
PRIOR FILING DATE: 1999-05-17
PRIOR FILING DATE: 1999-05-17
PRIOR FILING DATE: 1999-05-17
NUMBER OF SEQ ID NOS: 35
SEG ID NO 19
LENTH: A0
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1larity 100.0%; Pred. No. 1.5e-19;
Conservative 0; Mismatches 0;
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US-09-876-388-33
Sequence 33, Application US/09876388
Patent No. US20020049153A1
GENERAL INFORMATION:
APPLICANT: Bridon, Dominique P.
APPLICANT: L'Archevque, Benoit
APPLICANT: Ezrin, Alan M.
APPLICANT: Holmes, Darren L.
PRIOR APPLICATION NUMBER: 09/302,596
PRIOR FILING DATE: 1999-04-30
NUMBER OF SEQ ID NOS: 13
SOFWARE: PATENTIN Ver. 2.0
SEQ ID NO 7
LENGTH: 39
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ORGANISM: Artificial Sequence
                                                                                                                                           TYPE: PRT;
CRGANISM: Heloderma suspectum
US-09-953-0218-7
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St. Pierre, Serge
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Best Local Similarity
Matches 39; Conserv
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Patent No. US20020147131A1
GENERAL INFORMATION:
APPLICANT: COOLIGAGE THOMAS L.
APPLICANT: COLIGAGE THOMAS L.
TITLE OF INVENTION: Metabolic Intervention with GLP-1 to Improve the Function of Isch
TITLE OF INVENTION: Reperfused Skeletal Muscle Tissue
FILE REFERENCE: P03660US6
CURRENT APPLICATION NUMBER: US/09/953,021B
CURRENT FILING DATE: 2001-09-11
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APPLICANT: ENLERS, MARIO.

TITLE OF INVENTION: TREATMENT OF ACUTE CORONARY SYNDROME WITH GLP-1.

FILE REFERENCE: 089187/0395

CURRENT TILING DATE: 2001-10-22

PRIOR APPLICATION NUMBER: 09/859,804

PRIOR FILING DATE: 2001-05-18

PRIOR FILING DATE: 2001-05-18

PRIOR FILING DATE: 2001-05-19

PRIOR FILING DATE: 2001-05-19

NUMBER OF SEQ ID NOS: 13
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; O'THER INFORMATION: Description of Unknown Organism: Exendrin 3
US-09-982-978-7
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                                                                                                                                                                                                                                                                                                                 NAME/KEY: AMIDATION
LOCATION: (39)...(39)
OTHER INFORMATION: amidated Ser (Serinamide)
US-09-003-869-1
EARLIER APPLICATION NUMBER: US 60/055,404
EARLIER FILING DATE: 1997-08-08
EARLIER APPLICATION NUMBER: US 60/065,442
EARLIER FILING DATE: 1997-11-14
EARLIER APPLICATION NUMBER: US 60/066,029
EARLIER FILING DATE: 1997-11-14
NUMBER OF SEQ ID NOS: 188
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7, Application US/09982978 Patent No. US20020146405Al GENERAL INFORMATION:
                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Heloderma horridum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 7
LENGTH: 39
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ORGANISM: Unknown Organism
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RESULT 12
US-09-756-690A-14
US-09-756-690A-14
Sequence 14, Application US/09756690A
PUBLICATION NO. US20030036504A1
GENERAL INFORMATION:
APPLICANT: VOLTERWAN, ORVILLE G.
TITLE OF INVENTION: TRIGLYCERIDE LEVELS AND TREATMENT OF DYSLIPIDEMIA
TITLE OF INVENTION: TRIGLYCERIDE LEVELS AND TREATMENT OF DYSLIPIDEMIA
FILE REPERENCE: 249/124
CURRENT PRILING DATE: 2002-04-19
PRIOR PPLICATION NUMBER: 06/175,365
PRIOR FILING DATE: 2000-01-10
NUMBER OF SEQ ID NOS: 188
SOFTWARE: Patentin Ver 2.1
SEQ ID NO 14
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US-Sequence 14. Application US/09003869A

Patent No. US20020137666A1

GENERAL INFORMATION:

APPLICANT: BEDETEY. NIGEL ROBERT ARNOLD

APPLICANT: BEDETEY. KATHRYN S.

APPLICANT: BEREEKEY.

ITLE OF INVENTION: USE OF EXEMDINS AND AGONISTS THEREOF FOR TITLE OF INVENTION: THE REDUCTION OF FOOD INTAKE

TITLE OF INVENTION: THE REDUCTION OF FOOD INTAKE

TITLE OF INVENTION: THE REDUCTION OF FOOD INTAKE

TITLE OF INVENTION: USE OF EXEMDINS AND AGONISTS THEREOF FOR TITLE OF INVENTION: USE OF 0003, 869A

CURRENT APPLICATION NUMBER: US 60/034,905

EARLIER FILING DATE: 1997-01-07

EARLIER FILING DATE: 1997-01-07

EARLIER FILING DATE: 1997-11-14

EARLIER FILING DATE: 1997-11-14

NUMBER OF SEQ ID NOS: 188

SEQUEN NO 14

SEQ ID NO 14

LENGTH: 39
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98.1%; Score 204; DB 9;
Best Local Similarity 97.4%; Pred. No. 4.5e-19;
Matches 38; Conservative 0; Mismatches 1.
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   Pred. No. 1.5
Mismatches
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ORGANISM: Artificial Sequence
   100.08;
                               Conservative
      Local Similarity
les 39; Conserv
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US-09-876-388-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Xaa represents Lys(E-MPA)-NH2-5TFA and where "E" represents Epsilus-09-876-388-33
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OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Peptide
NAME/KEY: MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: Peptide NAME/KEY: MOD_RES LOCATION: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 40;
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APPLICANT: Ezrin, Alan M.
APPLICANT: Ezrin, Alan M.
APPLICANT: Bolme, Darren L.
APPLICANT: Holme, Darren L.
APPLICANT: Holme, Darren L.
APPLICANT: Holme, Darren L.
APPLICANT: St. Pierre, Serge.
FILE OF INVENTION: LONG LESTING INSULINOTROPIC PEPTIDES
FILE REPRENENCE: 500862010160
CURRENT APPLICATION NUMBER: 09/623,618
PRIOR FILING DATE: 2000-09-05
PRIOR PRILING DATE: 2000-09-05
PRIOR PRILING DATE: 1909-10-15
PRIOR FILING DATE: 1909-10-15
PRIOR FILING DATE: 1999-10-15
TITLE OF INVENTION: LONG LASTING INSULINOTROPIC PEPTIDES
FILE REFERENCE: 500862001610
CURRENT APPLICATION NUMBER: US/09/876,388
CURRENT FILING DATE: 2001-09-24
PRIOR PEPLICATION NUMBER: 09/623,618
PRIOR PEPLING DATE: 2000-09-05
PRIOR FILING DATE: 2000-09-05
PRIOR PEPLING DATE: 2000-09-05-17
PRIOR PEPLING DATE: 1999-10-15
PRIOR PEPLING DATE: 1999-10-15
PRIOR PEPLING DATE: 1999-10-15
PRIOR PEPLING DATE: 1999-05-17
NUMBER OF SEQ ID NOS: 35
SOFTWARE: PATENTIN VET: 2.1
SEQ ID NO 33
LENGTH: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 20%; DB 10; Best Local Similarity 100.0%; Pred. No. 1.5e-19; Matches 39; Conservative 0; Mismatches 0;
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APPLICANT: Bridon, Dominique P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
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LENGTH: 40
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DB 10; Length 40;

100.0%; Score 208;

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RESULT 14

U.S.-09-756-690A-2

Sequence 2, Application US/08756690A

Sequence 2, Application No. US20030036504A1

GENERAL INNORMATION:
APPLICANT: NOLTERAN, ORVILLE G.
TITLE OF INVENTION: TRIGITCERIDE LEVELS AND TREATMENT OF DYSLIPIDEMIA

FILE REFERENCE: 249/124

CURRENT APPLICATION NUMBER: US/09/756,690A

CURRENT APPLICATION NUMBER: 60/175,365

PRIOR FILING DATE: 2000-01-10

NUMBER OF SEQ ID NOS: 188

SOFTWARE: Patentin Ver 2.1

SEQ ID NO 2

LENGTH: 39

"LENGTH: 39

"THE OF INVENTION OF THE OF 
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| Sequence 12, Application US/09876388
| Patent N. US/0020049133A1
| GENERAL INFORMATION:
| APPLICANT: Bridon, Dominique P. APPLICANT: Extin, Alan M. APPLICANT: Extin, Alan M. APPLICANT: LATCHOVEQUE, BENOİK
| APPLICANT: LADIMES, DATEN L. APPLICANT: LADIMES, DATEN L. APPLICANT: LADIMES, DATEN L. APPLICANT: LADIMES, DATEN ST. PILE REFRENCE: 500862001610
| CURRENT APPLICATION NUMBER: US/09/876,388 |
| CURRENT APPLICATION NUMBER: 09/62,618 |
| PRIOR FILING DATE: 2000-09-05 |
| PRIOR FILING DATE: 2000-05-17 |
| PRIOR FILING DATE: 2000-05-17 |
| PRIOR FILING DATE: 1999-10-115 |
| PRIOR FILING DATE: 1999-10-115 |
| PRIOR FILING DATE: 1999-05-17 |
| PRIOR FILING DATE: 1999-05-17 |
                                                                                                                                                                                                                                                 Length 39;
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96.2%; Score 200; DB 9; Length 39
Best Local Similarity 94.9%; Pred. No. 1.4e-18;
Matches 37; Conservative 1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                     1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 HGDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
                                        NAME/KEY: AMIDATION

LOCATION: (39)...(39)

OTHER INFORMATION: amidated Ser (Serinamide)
US-09-003-869-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Heloderma suspectum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Exendin-4 FEATURE:
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US-09-876-388-12
FEATURE:
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NUMBER OF SEQ ID NOS: 35

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 12

LENGTH: 39

TYPE: PRT

ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: Deptide
US-09-876-388-12

Query Match
Best Local Similarity 94.9%; Pred. No. 1.4e-18;
Matches 37; Conservative 1; Mismatches 1; Indels 0; Gaps

OY 1 HSDGTFTSDLSKOWEEEAVELFIEWLKNGCPSSGAPPPS 39

I HSDGTFTSDLSKOWEEEAVELFIEWLKNGCPSSGAPPPS 39

Search completed: March 28, 2003, 08:41:09
Job time: 15 secs
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March 28, 2003, 08:35:18; Search time 43 Seconds (without alignments) 87.192 Million cell updates/sec
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                                                                                                                                                                                     US-09-756-690A-1
208
1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSŚGAPPPS 39
GenCore version 5.1.4_p5_4578
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                           283224 seqs, 96134422 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                        OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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1: pir1:*
2: pir2:*
3: pir3:*
1: pir4:*
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                                                                                                                                                                                                            Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Database :
                                                                                                                                                                                                                                                                                                                             Searched:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	1	exendin-4 - Gila m	precu	65	glucagon G1 - Nort	precu		pr	•	Пр	glucagon II precur		ı.	-1			glucagon precursor	gon	1	•		_	glucagon precursor	glucagon precursor	glucagon precursor	_	_	_	glucagon precursor
SUMMARIES	QI	HWGH32	HWGH4G	GCFGB	S44472	S44471	GCIDC	S44473	GCGXA	151093	151058	151057	GCAF2	B61125	C61125	GCDF	GCONC	вссн	151301	GCFLE	S07211	A61135	GCFIS	GCPG	ССНО	GCGP	GCRIDU	GCRT	GCHY	GCBO
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	Length	39	39	101	31	31	63	30	72	99	178	178	122	30	30	29	9	151	206	5	29	29	87	158	180	180	180	180	180	180
	- a :	0.0	?	49.0	9.6	9.7	9.7	7	7	4.7	4.7	4.7	8.	3.3	.3	2.3	2.3	L.3	1.3	6.0	6.0	6.0	6.0	6.0	6.0	6.0	6.0	6.0	6.0	6.0
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exendin-4 - Gila monster

RESULT 2

glucagon precursor glucagon l precurs glucagon - Chinchi glucagon - rabbit glucagon - common glucagon - common glucagon - burop glucagon - burop glucagon - turkey glucagon - turkey glucagon - cotrich glucagon - elephan glucagon - elephan glucagon - siider glucagon - siider glucagon - siider	ALIGNMENTS	HRESULT 1  HWGH3Z  Evandin-3 - Mexican beaded lizard  Evandin-3 - Mexican beaded lizard  C.Species: Heloderma horridum (Mexican beaded lizard)  C.Species: Heloderma horridum (Mexican 31-Mar-1993 #text_change 21-Nov-1997  C.Species: Heloderma horridum (Mexican 31-Mar-1993 #text_change 21-Nov-1997  C.Species: Heloderma horridum (Mexican 31-Mar-1993 #text_change 21-Nov-1997  C.Accession: A.Species: Loss
A57294 GCCB GCCB A91741 A91742 CG0840 GCOPV A91740	ALIG	Card (Mexican be revision 31 inman, W.A.; 62, 1990 ucture of ex UID:91056067 end; duplic carboxyl end carboxyl end: 0,0%; Score carboxyl end: 0,0%; Pred: 10,0%;
0011110101010110		lizard dum (Me: nce_rev nce_rev Xleinmal 20262, structu; MUID: MUID: 100.0%; 1ve EAVRLEI EAVRLEI EAVRLEI
1180 124 124 129 129 129 130 130 130		Mexican beaded lizard Heloderma horridum (Mexican beamar-1993 #sequence_revision 31- 18. A23674 Andrews, P.C.; Kleinman, W.A.; Andrews, P.C.; Kleinman, W.A.; Andrews, P.C.; Kleinman, W.A.; Arification and structure of exernitation and structure of exernitation and structure of exernitation and structure of exernitation of amylase.  17. 9 cENG. 17. 9 ucagon andated carboxyl end; duplicated site: andated carboxyl end ch is imilarity 100.0%; Score; 39; Conservative 0; MismalsDESKOMEEERVRLFIEWIKNGGEP
o4044440000000000000000000000000000000		a be
44 W W W W W W W W W W W W W W W W W W		Mexican beac eloderma hon far-1993 #sec A23674 an 265, 2021 an 265, 2021 number: A23 A23674 ype: proteil rification a number: A23 A2367 ype: proteil ype: proteil
68888888888888888888888888888888888888		HWGH3Z  exendin-3 - Mexican beaded lizard  exendin-3 - Mexican beaded lizard  c.Specis: Heloderma horridum (Mexican  c.Accession: A23674  B. Biol. Chem. 265, 20259-20262, 1990  A. Title: Purification and structure of  A. Recession: A23674  A. Molecule type: procein  A. Residues: 1-39 cENG  A. Molecule type: procein  A. Residues: 1-39 cENG  C. Comment: Exendins are venom component  g in secretion of amylase.  C. Superfamily: glucagon  C. Superfamily: glucagon  C. Skeywords: amidated carboxyl end; Mupl  F; 39/Modified site: amidated carboxyl  Guery Match  Best Local Similarity 100.0%; Scon  Best Local Similarity 100.0%; Pree  Matches 39; Conservative 0; Mi  I HSDGTFTSDLSKQMEEEAVRLFIEWIKN  Db  I HSDGTFTSDLSKQMEEEAVRLFIEWIKN  Db
0.00 0.00 0.00 0.00 0.00 0.00 0.00 0.0		RESULT 1 HWGH3Z exendin-3 C; Species 3 C; Date: 33 C; Accessing R; Eng. J. J. Blol. J. J

C. Species: Heloderma Suspectum (Gila monster)
C. Species: Heloderma Suspectum (Gila monster)
C. Date: 31-Mar-1993 #sequence\_revision 31-Mar-1993 #text\_change 21-Nov-1997
C. Date: 31-Mar-1993 #sequence\_revision 31-Mar-1993 #text\_change 21-Nov-1997
C. Date: 31-Mar-1993 #sequence\_revision 31-Mar-1993 #text\_change 21-Nov-1997
B. Bool. Chem. 267, 7400-7405, 1992
J. Biol. Chem. 267, 7400-7405, 1992
J. Marcel Isolation and characterization of exendin-4, an exendin-3 analogue, from B. A. Reference number: A42466; MUID:92218391; PMID:1313797
A. Reference number: A42466; MUID:92218391; PMID:1313797
A. Residues: protein
A. Residues: 1-39 < Comment: Exendin-4 does not stimulate amylase secretion by pancreatic acinar cell
C. Comment: Exendin-4 does not stimulate amylase secretion by pancreatic acinar cell
C. Superfamilly: glucagon
C. Reywords: amidated carboxyl end (Ser) #status experimental ö ; Length 39; Ouery Match 96.2%; Score 200; DB 1; Length 39 Best Local Similarity 94.9%; Pred. No. 4.1e-18; Matches 37; Conservative 1; Mismatches 1; Indels

1 HSDGTETSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39

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Gaps

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Oglucagon-like peptide - North American paddlefish (Polyodon spathula)
C;Species: Polyodon spathula
C;Species: S44473
R;Nguyen, T.M.; Mommsen, T.P.; Mims, S.M.; Conlon, J.M.
Biochem J. 300, 339-345, 1994
A;Nguyen, T.M.; Mommsen, T.P.; Mims, S.M.; Conlon, J.M.
A;Title: Characterization of insulins and proglucagon-derived peptides from a phyloge A;Reference number: S4467; MUID:94271144; PMID:8002937
A;Nccession: S4447
A;Molecule type: protein
C;Superfamily: glucagon
C;Superfamily: glucagon
C;Keywords: duplication; hormone; pancreas
F;1-30/Product: glucagon-like peptide #status predicted <MAT>
R;Nguyen, T.M.; Mommsen, T.P.; Mims, S.M.; Conlon, J.M.
Biochem. J. 300, 339-345, 1994
Birtle: Characterization of insulins and proglucagon-derived peptides from a phyloge
Birtle: Characterization of insulins and proglucagon-derived peptides from a phyloge
Birtle: Characterization of insulins and proglucagon
Birtle: Characterization of insulins and proglication; hormone; pancreas
C; Reywords: carbohydrate metabolism; duplication; hormone; pancreas
F;1-31/Product: glucagon G1 *status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Glücagon precursor - channel catfish (fragments)
C;Species: Ictalurus punctatus (channel catfish)
C;Species: Ictalurus punctatus (channel catfish)
C;Bacte: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 20-Mar-1998
C;Accession: A05166, A05167
R;Andrews, P.C.; Ronner, P.
J. Blol. Chan. 260, 3910-3914, 1985
J. Blol. Chan. 260, 3910-3914, 1985
A;Fitle: Isolation and structures of glucagon and glucagon-like peptide from catfish A;Reference number: A92514; MUID:85157336; PMID:3838546
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A,Residues: 30-63 <AND2>
C,Superfamily: glucagon
C,Keywords: carbohydrate metabolism; duplication; hormone; pancreas
F;1-29/Product: glucagon #status experimental <GCN>
F;30-63/Product: glucagon-like peptide 1 #status experimental <GLI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
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                                                                                                                                                                                                                                                                                                                                                                                                          Length 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             47.6%; Score 99; DB 1; Let 51.6%; Pred. No. 2.1e-05; Live 9; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46.2%; Score 96; DB 2; Le
58.6%; Pred. No. 2.2e-05;
tive 5; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                      Score 99; DB 2;
Pred. No. 9.9e-06;
5; Mismatches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      |:|||:|||:|||:|||:|| 30 HADGTYTSDVSSYLQDQAAKDFITWLKSGQP 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGP 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
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Best Local Similarity 58.6%;
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: protein A; Residues: 1-29 <AND1>
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Matches 16; Conserv
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Best Local S:
Matches 17,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                   NAlternate names: oxyntomodulin
N;Contaigs: glucagon; glucagon-36 (oxyntomodulin); glucagon-like peptide 1; glucagon-lik
C;Specides: Rana catesbeiana (bullifrog)
C;Date: i3-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 20-Mar-1998
C;Accession: B28091; C28091; D28091
R;Pollock, H.G.; Hamilton, J.W.; Rouse, J.B.; Ebner, K.E.; Rawitch, A.B.
A;Tille: Isolation of peptide hormones from the pancfeas of the bullfrog (Rana catesbeig
A;Reference number: A92730; MUID:88257102; PMID:3260236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ringuyen, T.M.; Mommsen, T.P.; Mims, S.M.; Conlon, J.M.
Blochem. J. 300, 339-345, 1994
Affitle: Characterization of insulins and proglucagon-derived peptides from a phylogenet A; Reference number: S44467; MUID:94271144; PMID:8002937
A. Accession: S44472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: protein
A; Residues: 1-31 < NGU>
A; Note: the sequence from Fig. 3 is inconsistent with that from Fig. 5 in having 29-Glu
C; Superfamily: glucagon
C; Superfamily: glucagon
C; Keywords: carbohydrate metabolism; duplication; hormone; pancreas
F;1-31/Product: glucagon G2 #status predicted < GCN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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C.Species: Polyodon spathula
C.Date: 18-Sep-1997 #sequence_revision 18-Sep-1997 #text_change 07-May-1999
C.Accession: S44471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             glucagon G2 - North American paddlefish (Polyodon spathula)
C;Species: Polyodon spathula
C;Date: 19-Mar-1997 #sequence_revision 12-Dec-1997 #text_change 07-May-1999
C;Accession: S44472
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Accession: D28091
A; Molecule type: protein
A; Molecule type: protein
A; Residues: 69-101 - P03>
C; Superfamily: glucagon
C; Superfamily: glucagon-36 (oxyntomodulin) #status experimental <G36>
P; 1-36/Product: glucagon #status predicted <GCN>
P; 37-67/Product: glucagon-like peptide 1 #status experimental <G11>
F; 37-67/Product: glucagon-like peptide 2 #status experimental <G11>
F; 69-101/Product: glucagon-like peptide 2 #status experimental <G11>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ouery Match
49.0%; Score 102; DB 1; Length 16
Best Local Similarity 54.8%; Pred. No. 1.5e-05;
Matches 17; Conservative 7; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 101; DB 2; L6
Pred. No. 5.6e-06;
6; Mismatches 6;
                            1 HGEGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGP 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
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Local Similarity 58.6%;
les 17; Conservative t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: protein A; Molecule type: protein A; Residues: 1.36 <P02> A; Accession: C28091 A; Molecule type: protein A; Molecule type: protein A; Residues: 37-68 <P0L>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S:
Matches 17
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djucagon II precursor - rainbow trout
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Species: Oncorhol: 151039, 151039, 151038
R) Irwin, Nong, J.
R) Areference number: A55895; MUD:95295739; PMID:776976
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Residues: 1-178 CIRW>
A;Residues: 1-178 CIRW>
A;Across-references: EMBL:U19914; NID:9736362; PIDN:AAC59668.1; PID:9736363
A;Title: Trout and chicken proglucagon: alternative splicing generates mRNA transcr
A;Reference number: A55895; MUID:95295739; PMID:7776976
A;Accession: IS108
A;Accession: IS108
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Cross.references: EMBL:U19917; NID:9736364; PIDN:AAC59669.1; PID:9736365; GB:S784
A;Accession: IS1299
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A; Residues: 'M', 114-144 < IR4>
A; Residues: 'M', 114-144 < IR4>
A; Cross-references: EMBL: U19919; NID: 9736374; PIDN: AAC60213.1; PID: 9736377
A; Accession: 151036
A; Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: DNA
A;Residues: 113-123 <IR5>
A;Cross references: EMBL:U19918; NID:g736373; PIDN:AAC60212.1; PID:g736376
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Molecule type: mRNA
A;Residues: 58-123 <IR3>
A;Cross-references: BMBL:U19913; NID:g736360; PIDN:AAC59667.1; PID:g736361
A;Status: preliminary; translated from GB/EWBL/DDBJ
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A; Residues: 113-144 CIR2>
A; Cross-references: EMBL:U19916; NID:g736369; PIDN:AAC60210.1; PID:g736372
A; Accession: 151038
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A;Residues: 113-123 <1R3>
A;Cross-references: EMBL:U19915; NID:q736368; PIDN:AAC60209.1; PID:g736371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                         A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 52-53, X', 55-123 <IR2>
A;Residues: 52-53, X', 55-123 <IR2>
A;Cross references: GB:S78473; NID:9999382; PIDN:AAB34504.1; PID:9999383
A;Accession: 151056
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     44.7%; Score 93; DB 2; 1
48.3%; Pred. No. 0.00036;
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44.7%; Score 93; DB-Best Local Similarity 48.3%; Pred. No. 0.00
Matches 14; Conservative 10; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Introns: 123/2
C; Superfamily: glucagon
C; Keywords: duplication
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A, introns: 123/2
C, Superfamily: glucagon
C, Keywords: duplication
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                                                                                                                                                                                                                                                       N:Conthins: glucagon; glucagon; glucagon; glucagon-like peptide 1
C;Species: Lepisosteus spatula (alligator gar)
C;Species: Lepisosteus spatula (alligator gar)
C;Date; 31-Mar-1993 segeunce_revision 31-Mar-1993 #text_change 20-Mar-1998
C;Date; 31-Mar-1993 segen
C;Accesion: S06339; S06871
R;Pollock, H.G.; Kimmel, J.R.; Ebner, K.E.; Hamilton, J.W.; Rouse, J.B.; Lance, V.; Rawi
Gen. Comp. Endocrinol. 69, 133-140, 1988
A;Reference number: S06339; MuID:88196798; PMID:3282974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Wolecule type: protein
A;Residues: 39-72 <PO2>
C;Comment: X's at positions 37-38 represent a pair of basic amino acids forming a cleava
C;Superfamily: glucagon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dlucagon - chinook salmon (fragment)
Cispecies: Oncorhypichus tschawytscha (chinook salmon)
Cispecies: Oncorhypichus tschawytscha (chinook salmon)
Cispecies: Oncorhypichus tschawytscha (chinook salmon)
Cispecies: Oncorhypichus tschawytscha 13-Sep-1996 #text_change 16-Jul-1999
Cisaccession: I51093
Ritthin, Dam, Wongy, J.
Moi. Endocrinol. 9, 267-277, 1995
A;Title: Trout and chicken proglucagon: alternative splicing generates mRNA transcripts
A;Accession: I51093
A;Acteus: predluminary: translated from GB/EMBL/DDBJ
A;Rotus: predluminary: translated from GB/EMBL/DDBJ
A;Rotus: predluminary: Aranslated from GB/EMBL/DDBJ
A;Rotus: predluminary: BMBL: U19920; NID:9736366; PIDN:AACS9670.1; PID:9736367
C;Superfamily: glucagon
C;Superfamily: glucagon
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C;Species: Oncorhynchus mykiss (rainbow trout)
C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 16-Jul-1999
C;Accession: 151058; 151299; 151056; 151037; 151036; 151300
R;Irwin, D.W.; Wong, J.
Mol. Endocrinol. 9, 267-277, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Keywords: carbohydrate metabolism; duplication; hormone; pancreas p.136/Pcroduct; glucagon-36 (cxyntcondoulin) #status experimental <G36>F:1-29/Product; glucagon #status predicted <GCN>F:39-70/Product; glucagon-like peptide l #status predicted <GL1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    / Match 45.2%; Score 94; DB 1; Length 72; Local Similarity 48.3%; Pred. No. 0.0001; nes 14; Conservative 9; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tch 44.7%; Score 93; DB 2; Length 66;
al Similarity 48.3%; Pred. No. 0.00012;
14; Conservative 10; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 66;
                                                                                                                                                                                                        glucagon precursor - alligator gar (fragment)
N.Alternate names: oxyntomodulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
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33 HADGTYTSDVSTYLQDQAAKDFVSWLKSG 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
          A.Molecule type: protein
A;Residues: 1-36 <POL>
A;Accession: S06871
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Matches
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IS1058
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                                                                                                                                           RESULT 8
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Gaps

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us-09-756-690a-1.rpr

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Jucagon - smaller spotted catshark
C;Species: Scyliorhinus canicula (smaller spotted catshark, smaller spotted dogfish)
C;Species: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_change 20-Mar-1998
C;Accession: A26992
FRSS Lett. 214, 50-56, 1987
A;Title: Primary structure of glucagon from the gut of the common dogfish (Scyliorhin A;Reference number: A26992, MuID:87190953; PMID:3569517
A;Title: The primary structure of glucagon-11ke peptide but not insulin has been cons A;Reference number: A61125; MUID:91340068; PMID:1874385 A;Accession: C61125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A,Molecule type: protein
A,Residues: 1-29 <CON>
C,Superfamily: glucagon
C,Keywords: carbohydrate metabolism; duplication; hormone; intestine; pancreas
                                                                                                                                                             A.Molecule type: protein
A.Molecule type: protein
A.Residues: 1730 <CON>
C.Superfamily: glucagon
C.Superfamily: glucagon
F.1-30/Products: amidated carboxyl end; duplication
F.1-30/Modified site: amidated carboxyl end (Arg) *status experimental
F.30/Modified site: amidated carboxyl end (Arg) *status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 88; DB 1; Length 29;
Pred. No. 0.00021;
5; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       43.3%; Score 90; DB 2; Length 30;
llarity 44.8%; Pred. No. 0.00012;
Conservative 10; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Search completed: March 28, 2003, 08:40:13
Job time : 43 secs
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Best Local Similarity 53.6%;
Matches 15; Conservative
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Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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C; Species: Anguilla rostrata (American eel)
C; Accession: B61125
R; Conlon, J.M.; Andrews, P.C.; Thim, L.; Moon, T.W.
Gen. Comp. Emdocrinol. 82, 23-32, 1991
A; Title: The primary structure of glucagon-like peptide but not insulin has been conservent ession: B61125
A; Reference number: A61125; MUID: 91340068; PMID: 1874385
A; Accession: B61125
A; Accessi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dincagon 2 precursor - American goosefish

Nicontains: glucagon; glucagon: like peptide 1

Nicontains: glucagon; glucagon: like peptide 1

C;species: Lobius americanus (American goosefish)
C;pate: 31-Mar-1993 #sequence_revision 31-Mar-1993 #text_change 21-Jul-2000
C;Accession: A05159
Biol. Chem. 258, 3280-3284, 1983
A;Fille: Anglerfish islet pre-proglucagon II. Nucleotide and corresponding amino acid & A;Reference number: A05150; MUD:83135785; PMID:6338015
A;Accession: A05150
A;Acc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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C;Species: Anguilla anguilla (European eel)
C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 21-Nov-1997
C;Accession: C61125
R;Conlon, J.M.; Andrews, P.C.; Thim, L.; Moon, T.W.
Gen. Comp. Endocrinol. 82, 23-32, 1991
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C;Keywords: carbohydrate metabolism; duplication; hormone; pancreas
C;Keywords: carbohydrate metabolism; duplication casto-
F:12-122/Product: proglucagon 2 #status predicted <PGC2>
F:52-80/Product: glucagon #status predicted <GCN>
F:89-119/Product: glucagon-11ke peptide 1 #status predicted <GLI>
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Local Similarity 48.3%; Pred. No. 0.00042;
hes 14; Conservative 9; Mismatches 6
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43.3%; Score 90; DB 2;
Best Local Similarity 44.8%; Pred. No. 0.00012;
Matches 13; Conservative 10; Mismatches 6
                                                       10; Mismatches
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                                                       Matches 14; Conservative
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C61125
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Gaps

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heloderma s canis famil sus scrofa homo sapien saccharomyc homo sapien rattus norv ovis aries cavia porce

09prq9 1 p09682 h p04203 h p012910 c p01282 h p15442 s p01583 h p11384 r p11384 r p11384 r p11384 r p11384 r

GLUC\_LAMFL
GLUC\_HYDCO
EXEL\_HELSU
SECR\_CAFFA
SECR\_CAFFA
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SECR\_HUMAN
SECR\_HUMAN
SECR\_RAT
SECR\_RAT
VIP\_CAVPO
VIP\_CAVPO

29 36 38 27 131 170 121 121 134 27

75 73 60 61 60 60 60 60 60 60

ALIGNMENTS

lampetra fl hydrolagus

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                  March 28, 2003, 08:25:52; Search time 25 Seconds (without alignments) 64.703 Million cell updates/sec
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208
1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSŚGAPPPS 39
GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                112892 seqs, 41476328 residues
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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NO.	Score	Match	Match Length	BB	QI	Description	tion
1	208	. 0	39	-	ехез_негно	P20394	heloderma h
~	200	96.2	87	Н	EXE4_HELSU	P26349	-
m	102	σ	71	٦	GLUC_ICTPU	P04093	ictalurus p
4	102	49.0	103	•	GLUC_RANCA		rana catesb
'n	86		71	•	GLUC_PIAME	P81880	plaractus m
ی ر	94	45.2	78		GLUC_LEPSP	P09566	lepisosteus
,	91	<u>س</u>	121	н	GLUC_CARAU	P79695	carassius a
. œ	91	43.	122		GLU2_LOPAM	P04092	
9	06	43	30	Н	GLUM_ANGAN	P41521	anguilla an
0	88	42.	59	٦	GLUC_SCYCA	P09687	scyliorhinu
1	88	42	89	Н	GLUC_ONCKI		oncorhynchu
17	86	4	151		GLUC_CHICK		gallus gall
13	82	40.	29	٦	GLUC_PLAFE		platichthys
14	85	40.	29	Н	GLUC_TORMA	P09567	torpedo mar
15	82	40,	96	-	GLUC_MYOSC	P09686	myoxocephal
16	82	40.	158		GLUC_PIG	P01274	
17	85	40.	180		GLUC_BOVIN	P01272	bos taurus
18	85	40.	180		GLUC_CAVPO	P05110	
19	85	40.	180		GLUC_HUMAN	P01275	homo sapien
20	85	40	180		GLUC_MESAU	P01273	mesocricetu
2	85	40.	180		GLUC_MOUSE	P55095	mns mnscnln
22	85	40,	180	٦	GLUC_OCTDE	P22890	octodon deg
23	8.5	40	180		GLUC_RAT	P06883	rattus norv
2 6	84	40	36		GLU1_ORENI	P81026	oreochromis
25	8	40.	124		GLU1_LOPAM	P01278	lophius ame
2 6	. 6	6	7		GLUC_CHIBR	P31297	chinchilla
2.5	0 00	30	2		GLUC_RABIT	P25449	oryctolagus
20	82	30	9	7	GLUC_CANFA	P29794	canis famil
6	49	80	е.		GLU2 ORENI	P81027	oreochromis
0	77	37.	~		GLUC_DIDMA	P18108	didelphis m
31	76	36			GLUC_ANAPL	P01276	anas platyr
32	76	36.	7	٦	GLUC_CALMI	P13189	lorhyn
33	76	36.	7	Т	GLUC_AMICA	P33528	amia calva

RESULT 1  PERSHELLO  EXESHELLO  EXESHELLO  PERS191 (Rel. 17, Created)  TO
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Siochem. Biophys. Res. Commun. 143:87-92(1987).
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                                                                                                                          Gaps
                           SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

Chen Y.E., Drucker D.J.;

Chen Y.E., Drucker D.J.;

"Itsaue-specific expression of unique mRNAs that encode proglucagon-
derayed peptides or exendin 4 in the lizard.";

plool. Chem. 272.4108-4115(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Glucagon precursor (Fragment).
Ictalurus punctatus (Channel Catfish).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostel; Ostariophysi; Siluriformes;
Ictaluridae; Ictalurus.
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AMIDATION (G-87 PROVIDE AMIDE GROUP).
656BA6E3D87454A2 CRC64;
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MEDLINE~87156787;
HODGLINE~87156787;
HODGLIN N.M., Mahrenholz A.M., Andrews P.C., Gurd R.S.;
"Biological activities of catfish glucagon and glucagon-like peptide.";
                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 200; DB 1; Length 87;
Pred. No. 5.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4 HGEGTFTSDLSKOMEEEAVRLFIEWLKNGGPSSGAPPPS 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
                                                                                                                                                                                                                                                                                                            EMBL, U77613; AABS1130.1; -.
PIR, A42486; HWGHG.
InterPro; IPR000552; Glucagon.
Pfam; PF00123; hormone2; 1.
SMART; SM00070; GLUCA; 1.
A PROSITE; PS00260; GLUCAGON; 1.
A Glucagon family; Toxin; Amidation; Signal.
A Glucagon family: Toxin; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-NOV-1986 (Rel. 03, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
01-NOV-1990 (Rel. 16, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     71 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                         96.2%; Score 200; DE ilarity 94.9%; Pred. No. 5.1e Conservative 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                   MEDLINE-92218391; Pubmed-1313797;
                                                                                                                                                                                                                                                                                                                                                                                                                      87 AA; 9479 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
tes 37; Conserv
                                                                                                SEQUENCE OF 48-86.
Heloderma.
NCBI_TaxID=8554;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=7998
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PO4093;
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SEQUENCE
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J. Biol. Chem. 260:3910-3914(1985).

J. Biol. Chem. 260:3910-3914(1985).

J. Biol. Chem. 260:3910-3914(1985).

THE BLOOD SUGAR LEVEL.

IN BESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.

IN RESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.

JAMERICANEDOUS: X'S IN THE SEQUENCE WERE INCLUDED BY HOWOLOGY WITH AMERICANEDOUS.

J. SIMILARITY: BELONGS TO THE GLUCAGON FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                Andrews P.C., Ronner P.; "Isolation and structures of glucagon and glucagon-like peptide from cafish pancreas.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GLUC_RANCA STANDARD; PRT; 103 ÅA. ; ; 1915.48; P15.48;                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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49.0%; Score 102; DB 1; Length 71;
Best Local Similarity 54.8%; Pred. No. 1.6e-06;
Matches 17; Conservative 8; Mismatches 6; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24688E79AD981A8F CRC64;
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GLUCAGON-LIKE PEPTIDE.
E -> D (IN REF. 2).
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MEDLINE-85157536; PubMed-3838546;
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MEDLINE-88257102; PubMed-3260236;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HSSP; P01274; IGCN.
InterPro: IPR000532; Glucagon.
Fdam; PF00123; hormone2; 2.
SMART; SM00070; GLUCA; 2.
PROSITE; PS00260; GLUCAGON; 2.
Glucagon family; Hormone.
NON TER 1 GLUCAGON; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HSSP; P01274; IGCN.
INICEPTO: IPRO00532; Glucagon.
PRIMTS; PR00775; GLUCAGON.
SMARY; SM00070; GLUCA; 3.
PROSITE; PS00260; GLUCAGON; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            71 AA; 8173 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=8400;
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CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE.
SECUENCE
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Insulin and pancreatic polypeptide.";

Gen. Comp. Endocrinol. 67:375-382(1987).

1- FUNCTION: PROMOTES HIDOLYSIS OF GLYCOGEN AND LIPIDS, AND RAISES

1- FUNCTION: PROMOTES HIDOLYSIS OF GLYCOGEN AND LIPIDS, AND RAISES

1- INDUCTION: PROMOTES IN THE A CELLS OF THE ISLETS OF LANGERHANS

IN RESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.

1- MISCELLANBOUS: X'S IN THE SEQUENCE WERE INCLUDED BY HOMOLOGY WITH

MISCELLANBOUS: X'S IN THE SEQUENCE WERE INCLUDED BY HOMOLOGY WITH

1- SIMILARITY: BELONGS TO THE GLUCAGON FAMILY.

1- SIMILARITY: ACCOUNTY OF THE GLUCAGON FAMILY.
01-NOV-1990 (Rel. 16, Last sequence update)
| Locy-2001 (Rel. 40, Last annotation update)
| Locy-2001 (Rel. 40, Last annotation update)
| Glucagon precursor [Contains: Glucagon; Glucagon-36 (Oxyntomodulin);
| Glucagon-like peptide] (Fragment).
| Clucagon-like peptide] (Fragment).
| Lepisosteus spatula (Alligator gar) (Atractosteus spatula).
| Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
| Actinopterygil; Neopterygil; Semionotiformes; Lepisosteidae;
                                                                                                                                                      TISSUE-Pancreas;
MEDLINE-88196798; PubMed-3282974;
MEDLINE-88196798; PubMed-3282974;
Pollock H.G., Kimmel J.R., Ebner K.E., Hamilton J.W., Rouse J.B.,
Lance V., Rawitch A.B.;
"Isolation of alligator gar (Lepisosteus spatula) glucagon,
oxyntomodulin, and glucagon-like peptide: amino acid sequences of
oxyntomodulin and glucagon-like peptide.";
Gen. Comp. Endocrinol. 69:133-140(1988).
                                                                                                                                                                                                                                                                                                                   Pollock H.G., Kimmel J.R., Hamilton J.W., Rouse J.B., Ebner K.E., Lance V., Rawitch A.B.; "Isolation and structures of alligator gar (Lepisosteus spatula)
                                                                                                                                                                                                                                                                                               TISSUE-Pancreas;
MEDLINE-88030594; PubMed-3311873;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00123; hormone2; 2.
SMART; SM00070; GLUCA; 2.
PROSITE; PS00260; GLUCAGON; 2.
GLUCAGON family; Hormone. GLUCAGON; 2.
PEPTIDE 1 36 GLUCAGON; 2.
PEPTIDE 45 GLUCAGON; 3.010; SEQUENCE 78 AA; 8990 MW; 3011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HSSP; P01274; 1GCN.
InterPro; IPR000532; Glucagon.
                                                                                                                                                                                                                                                                              PRELIMINARY SEQUENCE OF 1-29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    45.2%;
                                                                                                                                         SEQUENCE OF 1-36 AND 45-78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                              NCBI_TaxID-7917;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GLUC_CARAU
P79695;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                     Glucagon precursor (Fragment).
Plaractus mesopotamicus (Pacu).
Bukaryota: Metazoa: Chordata: Craniata; Vertebrata; Euteleostomi;
Actinopterygii: Neopterygii; Teleostei; Ostarlophysi; Characiformes; Characidae; Plaractus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
                                                                                                                                             ;
                                                                                                                 Length 103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         47.1%; Score 98; DB 1; Length 71; illarity 51.6%; Pred. No. 5.2e-06; Conservative 9; Mismatches 6; Indels
                                GLUCAGON-36 (OXYNTOMODULIN). GLUCAGON-LIKE PEPTIDE 1.
                                                                                                              49.0%; Score 102; DB 1; Length 10
11arity 54.8%; Pred. No. 2.4e-06;
Conservative 7; Mismatches 7; Indels
                                36 GLUCAGON-36 (OXYNTOMODULIN
70 GLUCAGON-LIKE PEPTIDE 1.
71 GLUCAGON-LIKE PEPTIDE 2.
11719 MW: 316287787BAREIC8F7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 71 AA; 8146 MW; F66A3CA2DD9806C5 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GLUCAGON.
GLUCAGON-LIKE PEPTIDE.
                                                                                                                                                                                                                                                                                                 30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ā
                                                                                                                                                                                                                                                                      71 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGP 31
                                                                                                                                                                        1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGP 31
                                                                                                                                                                                       GLUCAGON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                      PRT;
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MEDLINE-99259587; PubMed-10327603;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Interpro; IPP000532; Glucagon. Pfam; PF00123; hormone2; 2. PRINTS; PR00275; GLUCAGON. SMARY; SM0070; GLUCA; 2. PROSITE; PS00260; GLUCAGON; Glucagon family; Hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GLUC_LEPSP STANDARD; F
P09566;
01-MAR-1989 (Rel. 10, Created)
        Glucagon family; Hormone.
PEPTIDE 1 29
                                                                                                                                                                                                                                                                        STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
Matches 16; Conserv
                                                                                         103 AA;
                                                                                                                                 Local Similarity
les 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=42528;
                                                                                                                                                                                                                                                                     GLUC_PIAME
P81880;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NON_TER
SEQUENCE
                                  PEPTIDE
PEPTIDE
NON_CONS
PEPTIDE
SEQUENCE
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GLUC_LEPSP
ID GLUC_LI
AC P09566
DT 01-MAR
                                                                                                                                    Best Loca
Matches
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Meopterygii; Teleostei; Ostarlophysi; Cypriniformes; Cyprinidae; Carassius.
                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
116-OCT-2001 (Rel. 40, Last annotation update)
Glucagon precursor [Contains: Glicentin-related polypeptide (GRPP);
Glucagon; Glucagon-like peptide)
Carassius auratus (Goldfish).
                                                                                                                                                  ö
                                                                                                      Score 94; DB 1; Length 78; Pred. No.1,9e-05; Mismatches 6; Indels
                                  GLUCAGON-LIKE PEPTIDE.
30106496271594E0 CRC64;
                                                                                                                                                                                                                                                                                                                                                            121 AA.
                                                                                                                                                      9; Mismatches
                   GLUCAGON-36.
GLUCAGON.
                                                                                                                                                                                                1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
                                                                                                                                                                                                                                SEQUENCE FROM N.A.

Yuen T.T.H., Mok P.Y., Chow B.K.C.;
                                                                                                                                                                                                                                                                                                                                                              PRT;
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                                                                                                                                                     HSSP, POLZ74; JGCN.
HSSP, POLZ74; JGCN.
PLEAPRO; IPR000232; Glucagon.
PRAM: PP00113; hormone2; 2.
PRINTS: PR00275; GLUCAGON.
PRART; SM00700; GLUCAA; 2.
PROSITE; PS00260; GLUCAA; 2.
Glucagon family; Hormone; Cleavage on pair of basic residues; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Divor.1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
01-NOV-1995 (Rel. 32, Last annotation update)
Glucagon-like peptide (GLP)
Anguilla anguilla (European freshwater eel), and
Anguilla rostrata (American eel).
Eukaryota: Metazoa: Chordata: Graniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Anguilliformes; Anguillidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conlon J.M., Andrews P.C., Thim L., Moon T.W.; The primary structure of glucagon-like peptide but not insulin has been conserved between the American eel, Anguilla rostrata and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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                                                                                                                                                                                                                                                                                                                                                                                                                43.8%; Score 91; DB 1; Length 122;
48.3%; Pred. No. 7.7e-05;
tive 9; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                   GLICENTIN-RELATED POLYPEPTIDE GLUCAGON II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            43.3%; Score 90; DB 1; Length 30;
44.8%; Pred. No. 2.1e-05;
Live 10; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                            GLUCAGON-LIKE PEPTIDE II. 5140AC47EF915519 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOD_RÈS 30 30 AMIDATION.
SEQUENCE 30 AA; 3376 MW; 592DA5EABD6E49D0 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
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MEDLINE=91340068; PubMed=1874385;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14171 MW;
                                                                                                               EMBL; V00632; CAA23905.1; -. PIR; A05150; GCAF2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 48.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=7936, 7938;
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P41521;
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SEQUENCE
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PEPTIDE
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Matches
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GLUM_ANGAN
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01-NOV-1986 (Rel. 03, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
Glucagon II precursor (Contains: Glicentin-related polypeptide (GRPP);
Glucagon II; Glucagon-like peptide II].
Lophius americanus (American goosefish) (Anglerfish).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Acanthomorpha; Paracanthopterygii; Lophilformes; Lophildae; Lophilus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro: IPR000532; Glucagon.
Prami, Pro00123; hormone2; 2.
Prami, Pro00123; hormone2; 2.
SMART; SM00070; GLUCAGO.
SMART; SM00070; GLUCAGO.
Glucagon family: Hormone; Cleavage on pair of basic residues; Signal.
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
              Submitted (FEB-1997) to the EMBL/GenBank/DDBJ databases.
-1- FUNCTION: PROMOTES HYDROLYSIS OF GLYCOGEN AND LIPIDS, AND RAISES THE BLOOD SUGAR LEYEL.
-1- SIMILARITY: BELONGS TO THE GLUCAGON FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
MEDLINE-81357858; PubHed-6338015;
Lund P.K., Goodman R.H., Montminy M.R., Dee P.C., Habener J.F.;
Lund P.K., Goodman R.H., Montminy M.R., Dee P.C., Habener J.F.;
"Anglerfish islet pre-proglucagon II. Nucleotide and corresponding
amino acid sequence of the cDNR.";
J. Blol. Chem. 258:3280-3284(1983).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
0
                                                                                                                                                                                                                                                                                                                                                                                                              POTENTIAL.
GLICENTIN-RELATED POLYPEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        43.8%; Score 91; DB 1; Length 121;
45.2%; Pred. No. 7.6e-05;
Live 10; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GLUCAGON-LIKE PEPTIDE.
SC1D4BEC1D26B9C6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  122 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 45.2%; Pred. No. 7.66 14; Conservative 10; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     GLUCAGON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 AA; 13527 MW;
                                                                                                                                                                                                                                                           EMBL; U65528; AAB39563.1; -.
HSSP; P01274; IGCN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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SEQUENCE
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PEPTIDE
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RESULT 12
GLUC_CHICK
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                                                                                                                                                                                                                                                                                                                                          MEDLINE-81190953; Pubmed-356917;
A COLOID J.M., O'Toole L., Thim L.;
COLOID J.M., O'Toole L., Thim L.;
COLOID J.M., O'Toole L., Thim L.;
RT "Primary structure of glucagon from the gut of the common dogfish
RT (Scyliorhinus canicula).";
RT (Scyliorhinus canicula).";
REBS Lett. 214:80-56(1987).
- PUNCTION: PROMOTES HYDROLYSIS OF GLYCOGEN AND LIPIDS, AND RAISES
THE BLOOD SUGAR LEVEL.
- INUNCTION: PROMOTED IN THE A CELLS OF THE ISLETS OF LANGERHANS
- INUNCTION: PROMOTED IN THE A CELLS OF THE ISLETS OF LANGERHANS
- IN RESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.
- INUNCTION: PROMOTES TO THE GLUCAGON FAMILY.
- PIR; A26992; GCDF.
- RSP; PO1274; IGCN.
- RSP; PO1274; IGCN.
- RSP; PO1273; hormone-2; 1.
- PRINTS; PRO0275; GLUCAGON.
- ROAITES PSO0260; GLUCAGON.
- ROAITES PSO0260; GLUCAGON.
- ROAITES PSO0260; GLUCAGON.
- SEQUENCE 29 AA; 3529 MW; 6FA96392086F0226 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE-86214328; PubMed-3520699;
Plisetskaya E., Pollock H.G., Rouse J.B., Hamilton J.W., Kimmel J.R.,
Gorbman A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Isolation and structures of coho salmon (Oncorhynchus kisutch) glucagon and glucagon.like peptide."; glucagon and glucagon.like peptide."; Fegul. Pept. 14:57-67(1986).
-1. FUNCTION: PROMOTES HYDROLYSIS OF GLYCOGEN AND LIPIDS, AND RAISES THE BLOOD SUGAR LEVEL.
-1. FUNCTION: PRODUCED IN THE A CELLS OF THE ISLETS OF LANGERHANS IN RESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                   Glucagon.
Spyllorhinus canlcula (Spotted dogfish) (Spotted catshark).
Sukaryota, Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyllorhinidae; Scyllorhinus.
NCBL_TaxID-7830;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-ARR-1988 (Rel. 07, Created)
01-ARR-1988 (Rel. 07, Last sequence update)
01-NOV-1990 (Rel. 16, Last annotation update)
01-NOV-1990 (Rel. 16, Last annotation update)
01-NOV-1990 (Rel. 16, Last annotation update)
01-NOV-1990 (Rel. 16, Last annotation)
01-NOV-1990 (Rel. 16, Last annotation)
01-NOV-1990 (Rel. 18, Nepterygil; Teleostei; Euteleostei;
01-NOV-1990 (Rel. 18, Nepterygil; Salmoniformes; Salmonidae; Oncorhynchus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 88; DB 1; Length 29;
Pred. No. 3.6e-05;
5; Mismatches 8; Indels
                                                                                                                           01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
01-JAN-1990 (Rel. 13, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                68 AA.
                                                                                           29 AA.
 1 HAEGIYTSDVSSYLQDQAAKEFVSWLKTG 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 HSDGTFTSDLSKQMEEEAVRLFIEWLKN 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         42.3%;
Local Similarity 53.6%;
les 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                                  SEQUENCE.
TISSUE-Pancreas;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=8019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GLUC_ONCKI
                                                                                               SCYCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Loca)
Matches
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GLUC_ONCKI
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C. -i- MISCELLANEOUS: X'S IN THE SEQUENCE WERE INCLUDED BY HOMOLOGY WITH AMERICAN GOOSEFISH SEQUENCES.

-i- MISCELLANEOUS: GLN-14 IS A UNIQUE SUBSTITUTION FROM LEUCINE IN COTHER KNOWN GLUCAGON SEQUENCES AND GLUCAGON-LIKE PEPTIDES.

-i- SIMILARITY: BELONGS TO THE GLUCAGON FAMILY...

BIR; JPO103; GCONC...

R HSSP; PO1274; JGCN.

R HSSP; PO1274; JGCN.

R RSP; PO0123; Andrones.; 2.

R RAMART; SM00070; GLUCA; 2.

R RNART; SM00070; GLUCAGON; 2.

RWART; SM00070; GLUCAGON; 2.

RWART; SM00070; GLUCAGON; 2.

R ROSITE; PS00260; GLUCAGON; 2.

R PROFITER I J GLUCAGON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 15, Last sequence update)
01-AUG-1990 (Rel. 18, Last annotation update)
Glucagon precursor.
Gallus gallus (Chicken), and
Meleagris gallopavo (Common turkey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom1;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HORM. Metab. Res. 4:360-363(1972).

-i FUNCTION: PROMOTES HYDROLYSIS OF GLYCOGEN AND LIPIDS, AND RAISES

-i FUNCTION: PROMOTES HYDROLYSIS OF GLYCOGEN AND LIPIDS, AND RAISES

-i INDUCTION: PRODUCED IN THE A CELLS OF THE ISLETS OF LANGERHANS

IN RESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.

-i- MISCELLANBOUS: THE COMPOSITION OF TURKEY GLUCAGON APPEARS TO BE

IDENTICAL WITH CHICKEN.

-i- MISCELLANBOUS: CHICKEN PREPROGLUCAGON DOES NOT CONTAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Nucleotide sequence determination of chicken glucagon precursor cDNA. Chicken preproglucagon does not contain glucagon-like peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SPECIES-Chicken;
MEDLINE-76069271; PubMed-1194290;
POLLOCK H.G., KImmel J.R.;
Chicken glucagon. Isolation and amino acid sequence studies.";
J. Biol. Chem. 250:9377-9380(1975).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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BEDLINE-73074118; PubMed-4645932;
Markussen J., Frandsen E.K., Heding L.G., Sundby F.;
"Turkey glucagon: crystallization, amino acid composition and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SPECIES-Chicken; TISSUE-Pancreas;
MEDLINE-90249492; PubMed-2338135;
Hasegawa S., Terazono K., Nata K., Takada T., Yamamoto H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 88; DB 1; Length 68;
Pred. No. 9.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                        38 68 GLUCAGON-LIKE PEPTIDE.
68 68 68 A3; 7810 MW; 50D14214FF204C04 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                       GLUCAGON.
GLUCAGON-LIKE PEPTIDE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           151 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 44.8%; Pred. No. 9.6e
Matches 13; Conservative 11; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        |:|||:||:||:|| ::::| : |: |||:|| 38 HADGTYTSNVSTYLQDQAAKDFVSWLKSG 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPOSITION, AND SEQUENCE OF 55-83.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEBS Lett. 264:117-120(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9031, 9103;
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SEQUENCE OF 55-83.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GLUC_CHICK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Okamoto H.
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NON_TER
SEQUENCE
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STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=7788;
                                                                                                                                                                                                                                                                                               GLUC_TORMA
P09567;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              marmorata."
                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GLUC_MYOSC
ID GLUC_M
AC P09686
DT 01-MAR
DT 01-MAR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 15
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bloinformatics Institute. There are no restrictions on its use by non-profit institutions as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or entered an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                PROSITE; PS00260; GLUCAGON; 3. Glucagon family; Hormone; Cleavage on pair of basic residues; Signal; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SPECIES-T. Obesus; TISSUE-Pancreas;
MEDLINE-92009994; PubMed-1916209;
MEDLINE-92009994; PubMed-1916209;
Mavarro I., Gutierrez J., Caliacch J., Rivera J., Planas J.;
Mavarro I., Gutierrez J., Caliacch J., Rivera J., Planas J.;
Misolation and primary structure of glucagon from the endocrine pancreas of Thunnus obesus.";
Gen. Comp. Endocrinol. 83:227-232(1991).
Gen. Comp. Endocrinol. 83:227-232(1991).
THE BLOOD SUGAR LEVEL.
THE BLOOD SUGAR LEVEL.
                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                  AMIDATION (G-148 PROVIDE AMIDE GROUP).
B6C0D87536C0AEB5 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thunnus obesus (Bigeye tuna).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii: Neopterygii; Teleostei; Buteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;
Pleuronectoidei; Pleuronectidae; Platichthys.

NCBL_TAXID-8260, 8241;
                                                                                                                                                                                                                                                                                                                                                                                                 ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SPECIES-P. flesus;
MEDLINE-87219793; PubMed-3556313;
Conlon J.M., Davis M.S., Thim L.;
"Primary structure of insulin and glucagon from the flounder (Platichthys flesus).";
Gen. Comp. Endocrinol. 66:203-209(1987).
                                                                                                                                                                                                                                                                                                                                                                        41.3%; Score 86; DB 1; Length 151;
48.3%; Pred. No. 0.00043;
tive 8; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                         GLUCAGON-LIKE PEPTIDE
             -1- SIMILARITY: BELONGS TO THE GLUCAGON FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1991 (Rel. 20, Created)
01-NOV-1991 (Rel. 20, Last sequence update)
01-JUN-1994 (Rel. 29, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Platichthys flesus (European flounder), and
                                                                                                                                                                                                                                                                                     PROGLUCAGON.
                                                                                                                                                                                                                                                                                                                                                                                                                                       1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
                                                                                                                                                                                                                                                                                                GLUCAGON
                                                                                                                                                                    HISSP, PO1274; IGCN.
InterPro; IPR000552; Glucagon.
Pfam: PF00123; hormone2; 2.
PRINTS; PR00275; GLUCAGON.
SWART; SW00070; GLUCA; 2.
                                                                                                                                                                                                                                                                                                                                                  17520 MW;
 GLUCAGON-LIKE PEPTIDE II
                                                                                                                                    EMBL; Y07539; CAA68827.1; -.
                                                                                                                                                                                                                                                                                                                                                                                     Sest Local Similarity 48.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
                                                                                                                                                                                                                                                                         22
151
83
118
147
                                                                                                                                                  PIR; S09992; GCCH.
PIR; A91740; A91740.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GLUC_PLAFE
P23062;
                                                                                                                                                                                                                                                                                                                        PEPTIDE
MOD_RES
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
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PROPEP
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                                                                                                                                                                                                                                                                           SIGNAL
                                                                                                                                                                                                                                                                                     CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gen. Comp. Endocrinol. 60:398-405(1985).
-!- FUNCTION: PROMOTES HYDROLYSIS OF GLYCOGEN AND LIPIDS, AND RAISES THE BLOOD SUGAR LEVEL.
-!- INDUCTION: PRODUCED IN THE A CELLS OF THE ISLETS OF LANGERHANS IN RESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.
-!- SIMILARITY: BELONGS TO THE GLUCAGON FAMILY.
PIR: S07211; 507211.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conlon J.M., Thim L.; "Primary structure of glucagon from an elasmobranchian fish. Torpedo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Torpedo marmorata (Marbled electric ray).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Squalea; Hypnosqualea; Pristiorajea; Batoidea;
Torpediniformes; Torpedinoidei; Torpedinidae; Torpedo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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Pred. No. 8.8e-05;
6; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                         Score 85; DB 1; Length 29;
Pred. No. 8.8e-05;
7; Mismatches 7; Indels
IN RESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION --- SIMILARITY: BELONGS TO THE GLUCAGON FAMILY.
PIR; 809348; GCFLE.
PIR; A61135; A61135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29 AA; 3511 MW; 04D96392086F0227 CRC64;
                                                                                                                                                                                                                                                                                       3508 MW; 77D5943208662E52 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
01-MAR-1989 (Rel. 10, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                              7; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 HSEGTFTSDYSKYLDNRRAKDFVQWLMN 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 HSDGTFTSDLSKQMEEEAVRLFIEWLKN 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 HSEGTFSNDYSKYLETRRAQDFVQWLKN 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE-Pancreas;
MEDLINE-86083105; PubMed-4076759;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Interpro; IPP000552; Glucagon.
Pfam: PP00123; hormone2; 1.
PRINTS; PR00275; GLUCAGON.
SMART; SM00070; GLUCA, 1.
PROSITE; PS00250; GLUCAGON; 1.
GLUCAGON family; Hormone.
SEQUENCE 29 AA; 3511 MW; 0408
                                                                                                                                                                                                                                    PROSITE; PS00260; GLUCAGON; 1. Glucagon family; Hormone. SEQUENCE 29 AA; 3508 MW; 77
                                                                                                       HSSP, P01274; IGCN.
InterPro; IPR000532; Glucagon.
Pfam, PP00123; hormone2; 1.
PRINTS; PR00275; GLUCAGON.
SMART; SM00070; GLUCA; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    40.9%;
50.0%;
                                                                                                                                                                                                                                                                                                                                              40.9%;
50.0%;
                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 50.09
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14; Conservative
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DT 16-CCT-2001 (Rel. 40, Last annotation update)

C clucagon precursor (Contains: Gillcentin-related polypeptide (GRPP);

SG Ulucagon, Glucagon-like peptide] (Fragment).

CS Myoxocephalus scorpius (Shorthborn sculpin) (Daddy sculpin)

C canthomorpha; Acanthopterygii, Nectabraris Euteleostei;

Actinopterygii, Neopterygii, Teleostei; Euteleostei; Nectabracis;

C Acathomorpha; Acanthopterygii, Percomorpha; Scorpaeniformes;

C CC CACCOSTINE—STORMENT (STORTH CONTROL STORTHON SCORPANIE).

RM 131.

RM 141.

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1 HSDGTFTSDLSKQMEEEAVRLFIEWLKN 28

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OM protein - protein search, using sw model	<pre>March 28, 2003, 08:33:48 ; Search time 85 Seconds (without alignments) 94.539 Million cell updates/sec</pre>	US-09-756-690A-1	1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPSSGAPPPS 39
earch, using	28, 2003, 00	756-690A-1	TFTSDLSKQME
- protein s	March		
OM protein	Run on	Title:	Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

671580 Total number of hits satisfying chosen parameters: 671580 seqs, 206047115 residues Searched:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SpTREMBL\_21:\*
1: sp\_archea:\*
2: sp\_bacteria:\*
3: sp\_fungi:\*
4: sp\_human:\*
5: sp\_invertebrate:\* sp\_virus:\*
sp\_vertebrate:\*
sp\_unclassified:\* sp\_rvirus:\*
sp\_bacteriap:\*
sp\_archeap:\* sp\_mammal:\* sp\_mhc:\* sp\_organelle:\* sp\_plant:\* sp\_rodent:\* Database :

Pred. No. 1s the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	042143 xenopus lae 041144 xenopus lae 041144 xenopus lae 081419 hoplobatrac 081409 oncorhynchu 091189 oncorhynchu 081189 orlorthinu 081191 petromyzon 012956 heloderma s 091410 gallus gall 085190 cants famil 089100 petromyzon 090400 petromyzon 090404 gallus petromyzon 090404 ambloplites 092xx3 rhizobium m
SUMMARIES	042143 042144 042144 091409 091971 091971 091189 091189 091180 091410 091410 091410 091410 09143 09143
DB	
a Query Match Length DB	266 219 220 220 220 178 178 180 180 120 120 121 120 131 131
% Query Match	0.004444444444444444444444444444444444
Score	1117 1009 933 87.55 87.55 87.65 87.65 87.65 87.65 87.65 87.65
Result No.	2 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1

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3 Q9USQB 10 Q8S3P9 2 Q9K5S1 2 Q9K5S1 2 Q9KD20 10 Q9KD20 10 Q9KE7 10 Q9KE7 11 Q9D2Z7 10 Q9KE7 11 Q9D2Z7 12 Q9URE7 13 Q9U94 13 Q9U94 14 Q9U94 16 Q9SXC1 17 Q9SSC1 18 Q9SSC1 19 Q9SSC1 10 Q9SSC1	
233 6614 2008 2008 2008 600 1010 1010 1010 1010 1010 1010 10	132 309 310
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1111 12222222222222222222222222222222	4 4 4 4 2 6 4 3

## ALIGNMENTS

	266 AA.		0 t a p 2 t a	ice update)	on: glucadon-like peptide 1A	-1B); glucagon-like peptide 1C	(GLP-1C); glucagon-like peptide 2 (GLP-2)].		Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,	Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;			CRICE	TCTMG:		Irwin D.M. Satkunaralah M. Wen Y., Brubaker P.L., Pederson R.A.,		lovel GLP-1-like peptides with	insulinotropic properties.";	920(1997).	YCOGEN AND LIPIDS, AND RAISES	ALTERNATIVE PRODUCTS: 2 ISOFORMS: 1 (SHOWN HERE) AND 2: ARE		I FAMILY.					-		Glucadon family; Hormone; Signal; Cleavage on pair of basic residues;	•
	PRT: 26		Created)	Last annota	ns: Glucado	ide 1B (GLP	ide 2 (GLP-	ed frog).	; Craniata;	Mesobatrach			IOO GILL BANG	ANALLYE SFL	3287:	Wen Y., Br		e encodes n		. 94:7915-7	LYSIS OF GL	I SOFORMS: 1	SPLICING.	HE GLUCAGON			u			Ľ	g. qnal; Cleav	e splicing.
	PRELIMINARY:		(TrEMBLrel. 05, Created)	(TremBirel 17.	precursor [Contai	lucagon-like pept	lucagon-like pept	Xenopus laevis (African clawed frog).	Metazoa; Chordata	trachla; Anura;	Xenopus.	3355;	and the contract of the contra	DEMOMBLE FROM N.D., AND ALLEANALLYE SPLICING.	MEDI,INE-97368292: PubMed-9223287:	Satkuparalah M.		proglucagon gen	oic properties.";	Acad. Sci. U.S.A	FUNCTION: PROMOTES HYDRO	LIVE PRODUCTS: 2	BY ALTERNATIVE SPLICING.	ITY: BELONGS TO THE GLUCAGON FAMILY	EMBL; AF004432; AAB65660.1;	1; 1GCN.		Pfam; PF00123; hormone2; 5.	PKINIS; PROOZOS GLUCAGON.	SMAKI; SMUOVIO; GLUCA; 5. PROSITE: PSAA26A: GLUCAGON: 5	nily; Hormone; Si	Multigene family, Alternative splicing.
LT 1	042143	042143;	01-JAN-1998	01 - JIIN - 2001	Glucadon I r	(GLP-1A); gl	(GLP-1C); gl	Xenopus laev	Eukaryota; N	Amphibia; Ba	Xenopodinae; Xenopus.	NCBI_TaxID=8355;	LT]	TISSHEDDANDERS.	MEDI, INE = 9736	Irwin D.M.	Wheeler M.B.;	"The Xenopus	insulinotrop	Proc. Natl.	-1- FUNCTION	-1- ALTERNAT	PRODUCED BY	-!- SIMILARITY:	EMBL; AF0044	HSSP; P01274; 1GCN.	InterPro; IF	Pfam; PF0012	PRINTS; PROC	DECETTE: DECE	Glucadon fam	Multigene fa
RESULT	1	AC.	占	3 5	E	E	DE	SO	႘	႘	ပ္ပ	ŏ	2 0	ָ עַ עַ	2 2	RA	RA	RT	RT	RL I	ပ္ပင်	3 5	ខ្ល	ပ္ပ	DR	DR.	DK.	DR t	Z t	ž č	X E	W F

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Eukaryota; Metazoā; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae;
01-WAR-2002 (TrEMBLrel. 20, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                        Hoplobatrachus rugulosus
                                                                                                                                                                                                                                       NCBI_TaxID=110072;
                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                             Hoplobatrachus.
                                                                                                  Proglucagon
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            DDT REPRESENTATION OF STREET BY STRE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfant, PF00123; hormone2; 4.
PRINTS; PR00775; GLUCAGON.
SMRRT; SM0070; GLUCA, 4.
SMOSTITE; PS00260; GLUCAN, 3.
Glucagon family; Hormone; Signal; Cleavage on pair of basic residues;
                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JAN-1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
61ucagon II precursor (Contains: Glucagon; glucagon-like peptide lA (GLP-1A); glucagon-like peptide lB (GLP-1B); glucagon-like peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Irwin D.M., Satkunarajah M., Wen Y., Brubaker P.L., Pederson R.A., Wheeler M.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Xenopus laevis (African clawed frog).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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                                                                                                                                                                                                                                             ch 56.2%; Score 117; DB 13; Length 266; 1 Similarity 59.4%; Pred. No. 1.5e-07; 19; Conservative 10; Mismatches 3; Indels
                                      GLOCAGON-LIKE PEPTIDE IA.
GLOCAGON-LIKE PEPTIDE IB.
GLOCAGON-LIKE PEPTIDE IC.
GLOCAGON-LIKE PEPTIDE IC.
MISSING (IN ISOFORM 2).
W; 544F7BBC20AF872C CRC64;
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142 173 GLUCAGON-LIKE PEPTIDE 1B.
180 211 GLUCAGON-LIKE PEPTIDE 1C.
219 AA; 25271 MW; ACC699233G36ZCEO CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             219 AA
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                                                                                                                                                                                                                                                                                                                                                                                                        1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNGGPS 32
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                      GLUCAGON
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                                                                            142 173 GLU
180 211 GLU
227 259 GLU
214 261 MIS
266 AA; 30951 MW; 5
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Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Xenopodinae; Xenopus.
NCBI_TaxID=8355;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Multigene family.
SIGNAL 1
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SEQUENCE
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Q8UWL9
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Yeung C. M., Chow B.K.C.;
"Identification of a proglucagon cDNA from Rana tigrina rugulosa that encodes two GLP-18."
"ITENTIFICATION OF a proglucagon cDNA from Rana tigrina rugulosa that encodes two GLP-18."
"INTERPOS TRY 18000532, Glucagon."
"INTERPOS TRY 18000532, Glucagon."
PRINTS; PRO0123; hormone2; 4.
PRINTS; PRO01070; GLUCAGON.
"SMART; SMO0070; GLUCAGON.
"SNART; SMO0060; GLUCAGON.
"SROUENCE 220 AA; 25615 MW; C72D926E7F89E381 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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MEDLINE-95295739; PubMed-7776976;
ITWIN D.M., Wong J.;
ITWIN C.M., WONG J.;
INCORDINATION C.M.,
INCORDINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oncorhynchus tschawytscha (Chinook salmon) (King salmon).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygli, Neopterygli; Teleostei; Euteleostei;
Protacanthopterygli; Salmoniformes; Salmonidae; Oncorhynchus.
NCBI_TaxID=74940;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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47.6%; Score 99; DB 13; Length 220;
Best Local Similarity 44.4%; Pred. No. 3.1e-05;
Matches 16; Conservative 9; Mismatches 11; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 72;
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SEQUENCE 72 AA; 8293 MW; 8584352B1C260A31 CRC64;
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Q91409; Q91232;
Q1-NOV-1996 (TEMBLrel. 01, Created)
O1-NOV-1996 (TEMBLrel. 01, Last sequence update)
O1-DEC-2001 (TEMBLrel. 19, Last annotation update)
PROGLUCAGON (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 HSDGTFTSDLSKOMEEEAVRLFIEWLKNGGPSSGAP 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
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|HADGIYISDVSIYLQDQAAKDFVSWLKSG 67
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RESULT

220 AA

PRT;

PRELIMINARY;

QBUWL9;

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PROSITE; PS00260; GLUCAGON; 2.
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SEQUENCE
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PEPTIDE
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Matches
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                                                                                                                                                                                                                                                                                 PEGUENCE FROM N.A., AND ALTERNATIVE SPLICING.
TISSUE-DISTAL SMALL INTESTINE, AND PANCREAS;
X TISSUE-DISTAL SMALL INTESTINE, AND PANCREAS;
XA INTENDEDING-95295739; PubMed-7776976;
XA ITWIN D.M., Wong J.,
ITWIN D.M., Wong J.,
ITWIN D.M., Wong J.,
ITWIN D.M., Wong J.,
ITWIN D.M., Wong J.,
ITWIN D.M., Wong J.,
ITWIN D.M., Wong J.,
ITWIN D.M., Wong J.,
ITWIN D.M., Wong J.,
ITWIN THE CALL SMALL STATE OF THE STATE OF THE BLOOD SUGAR LEVEL (BY SIMILARITY).

THE BLOOD SUGAR LEVEL (BY SIMILARITY).

THE BLOOD SUGAR LEVEL OF YEAR PRODUCED BY ALTERNATIVE SPLICING.

INDICTION: PRODUCED IN THE A CELLS OF THE ISLETS OF LANGERHANS IN RESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.

INDICTION: PRODUCED IN BLOOD SUGAR CONCENTRATION.

INDICTION: ARCSPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.

INDICTION: ARCSPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.

INDICTION: ARCSPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.

EMBL: U19913; AAC650212.1; ...

EMBL: U19918; AAC660212.1; ...

EMBL: U19918; AAC60212.1; ...

EMBL: S78475; AAB34508.1; ...

EMBL: S78475; ABB34508.1; ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE: PSO0260; GLUCAGON; 3.
Slucagon family; Hormone; Cleavage on pair of basic residues; Signal;
Alternative splicing; Multigene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GRPP (GLICENTINE RELATED POLYPEPTIDE). GLUCAGON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 17, Last sequence update)
01-UND-2001 (TrEMBLrel. 17, Last annotation update)
Glucagon II precursor.
Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Eukaryotas Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygli; Neopterygli; Teleostei; Euteleostei;
Protacanthopterygli; Salmoniformes; Salmonidae; Oncorhynchus.
                                                                                                                                Glucagon I precursor.

Ohcorhynchus mykiss (Rainbow trout) (Salmo gairdneri).

phearyota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygil; Neopterygil; Teleostel; Buteleostel;

protacanthopterygil; Salmoniformes; Salmonidae; Oncorhynchus.

NCBI_TaxiD=8022;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GLUCAGON-LIKE PEPTIDE 1.
GLUCAGON-LIKE PEPTIDE 2.
MISSING (IN PANCREATIC ISOFORM).
5. SCR6980CF2A9D58E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  44.7%; Score 93; DB 13; Length 178;
11arity 48.3%; Pred. No. 0.00016;
Conservative 10; Mismatches 5; Indels
                                         091971; 091408; 091188; 092169;
01-NOV-1996 (TYEMBLEAL 01, Greated)
01-NOV-1996 (TYEMBLEAL 01, Last sequence update)
01-JUN-2001 (TYEMBLEAL 17, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT; 178 AA.
                      PRT; 178 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    52 80 GLC
85 120 GLC
137 169 GLC
124 178 MIS
178 AA; 20034 MW; 5
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Pfam; PF00123; hormone2; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRINTS; PR00275; GLUCAGO
SMART; SM00070; GLUCA; 3
                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Glucagon family; Hormo
Alternative splicing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
wes 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          091189; 092168;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HSSP; P01274;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VARSPLIC
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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Matches
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PH 10 1180
PH 
091971
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REPORTION N.A., AND ALTERNATIVE SPLICING.

TISSUE-DISTAL SWALL INTESTINE, AND PANCREAS;

RX MEDLINE-92295739; PubMed=7776976;

RT Trout and chicken proglucagon: alternative splicing generates mRNA irranscripts encoding glucagon-like peptide 2.";

MOI. Endocrinol. 9:267-277(1995).

-I- FUNCTION: PROMOTES HYDROLYSIS OF GLXCOGEN AND LIPIDS, AND RAISES THE BLOOD SUGAR LEVEL (BY SIMILARITY: BLOOD SUGAR LEVEL (BY SIMILARITY: SPLICING.)

-I- FUNCTION: PRODUCTS: 2 ISSOFORMS: INTESTINAL (SHOWN HERE) AND PANCREATIC; ARE PRODUCED BY ALTERNATIVE SPLICING.

-I- ALTERNATIVE PRODUCED BY ALTERNATIVE SPLICING.

-I- SIMILARITY: BELONGS TO THE GLUCAGON FAMILY.

DR RESPONSE TO A DROP IN BLOOD SUGAR CONCENTRATION.

-I- SIMILARITY: BELONGS TO THE GLUCAGON FAMILY.

DR REMEL; 019915; AAC60210:1; -..

DR REMEL; 019915; AAC60210:1; -..

DR REMEL; 019915; AAC60209:1; -..

DR REMEL; 019915; AAC60209:1; -..

DR RINC: 109015; ACCOMPANIENT ALTONOMEN.

DR RINC: 100013; CHUCAGON.

DR RINCE: DROUGTS: ALTONOMEN.

DR REMEL; DROUGTS: CHUCAGON.

DR RAILY: DROUGTS: CHUCAGON.

DR
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101-MAX-2000 (TERBLEAL 16, Last sequence update)
01-MAX-2000 (TERBLEAL 16, Last sequence update)
01-JUN-2002 (TERBLEAL 12, Last annotation update)
01-JUN-2002 (TERMELAL 12, Last annotation update)
Glucagon precursor (Contains: glucagon-29; glucagon-13; glucagon-11ke peptide) (Fragments).
Scyliorhius canicula (Spotted dogfish) (Spotted catshark).
Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes; Blasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes; Scyliorhinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pair of basic residues; Signal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GRPP (GLICENTINE RELATED POLYPEPTIDE) GLUCAGON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GLUCAGON-LIKE PEPTIDE 1.
GLUCAGON-LIKE PEPTIDE 2.
GLUCAGON-LIKE PEPTIDE 2.
GLUCAGON-LIKE PEPTIDE 2.
F. E89D73866CD91C66 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               44.7%; Score 93; DB 13; Length 178; tlarity 48.3%; Pred. No. 0.00016; Conservative 10; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      62 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSITE; PS00260; GLUCAGON; UNKNOWN_2. Glucagon family; Hormone; Cleavage on p Alternative splicing; Multigene family. 7 POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |:|||:|||:|
90 HADGTYTSDVSTYLQDQAAKDFVSWLKSG 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 HSDGTFTSDLSKQMEEEAVRLFIEWLKNG 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              178 AA; 19998 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
es 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE-PANCREAS;
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012956
ID 012956
AC 012956;
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             ä
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-1- FUNCTION: PROMOTES HYDROLYSIS OF GLYCOGEN AND LIPIDS, AND RAISES THE BLOOD SUGAR LEVEL.
-1- SIMILARITY: BELOONS TO THE GLUCAGON FAMILY.
EMBL, AF159707; AAF09186.1; -.
EMBL, AF159707; AAF09186.1; -.
InterPro: IPRO00532; Glucagon.
Pfam; PF00123; hormone2; 2.
PRINTS; PRO0275; GLUCAGON.
SWART; SM00070; GLUCAA. 2.
PROSTIE; PS00260; GLUCAGON. 2.
PROSTIE; PS00260; GLUCAGON. 2.
PROSTIE; PS00260; GLUCAGON. 2.
                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE-INTESTINE;

MEDILINE-940101712; pubmed-8405897;

GOALON J.M., Nielsen P.F., Youson J.H.;

"Primary structures of glucagon and glucagon-like peptide isolated from the intestine of the parasitic phase lamprey Petromyzon maxinus";
                                                                                                                                                                                                  OSPURI, OSPRAT,

101-MAY-2000 (TERMELRE). 13, Created)

101-MAY-2000 (TERMELRE). 13, Last sequence update)

101-MAY-2000 (TERMELRE). 13, Last sequence update)

101-DE-2001 (TERMELRE). 19, Last annotation update)

Glucagon I precursor (Contains: Glucagon; glucagon: like peptide 1 (GLP-1); glucagon. Itke peptide 2 (GLP-2)].

Everromyzon marinus (Sea lamprey).

Everromyzon (Sea lamprey).

Everromyzontiformes; Petromyzontidae; Petromyzon.
                                                                                                                                                                                                                                                                                                                                                         MEDLINE-20022986; PubMed-10555286;
Irwin D.M.; Huner O., Youson J.H.;
"Lamprey proglucagon and the origin of glucagon-like peptides.";
Mol. Blob: Evol. 16:1548-1557(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3;
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Pred. No. 0.00077;
5; Mismatches 8; Indels 3;
                                                                          Score 87.5; DB 13; Length 62; Pred. No. 0.00026;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GLUCAGON-LIKE PEPTIDE 1.
GLUCAGON-LIKE PEPTIDE 2.
9952C530D5A74072 CRC64;
                                           GLUCAGON-LIKE PEPTIDE.
C5FF487C12C69CD1 CRC64;
                                                                                                                                   '1 HSDGTFTSDLSKQMEEEAVRLFIEWL---KNGGPSSG 34
                                                                                       Pred. No. 0.000
6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 HSDGTFTSDLSKQMEEEAVRLFIEWL---KNGG 30
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GLUCAGON-33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           POTENTIAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          130 160 G
160 AA; 18042 MW;
            29
33
34
62
7270 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE OF 43-71 AND 82-113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 51.5%;
Matches 17; Conservative
                                                                          Match 42.1%;
focal Similarity 45.9%;
es 17; Conservative
 Glucagon family; Hormone
                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                     62 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Multigene family.
                                                                                                                                                                                                                                                                                                                                                TISSUE-INTESTINE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PEPTIDE
PEPTIDE
SEQUENCE
                                 NON_CONS
PEPTIDE
                                                      SEQUENCE
                                                                           Query Match
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                                                                                       Best Loc
Matches
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RESULT 9

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GRPP (GICENTINE RELATED POLYPEPTIDE).

GLUCAGON.-LIKE PEPTIDE 1.

GLUCAGON-LIKE PEPTIDE 2.

GLUCAGON-LIKE PEPTIDE 2.

D.> E (IN ISOPORM LPI).

MISSING (IN ISOFORM LPI).

MISSING (IN ISOFORM LPI).

MISSING (IN ISOFORM LPI).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauría; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                              Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Lepidosauria: Squamata, Scleroglossa; Anguimorpha, Helodermatidae;
                                                                                                                                                           SEQUENCE FROM N.A., ALTERNATIVE SPLICING, AND TISSUE SPECIFICITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           41.8%; Score 87; DB 13; Length 204; 51.7%; Pred. No. 0.0012; tive 6; Mismatches 8; Indels
                                          Last sequence update)
Last annotation update)
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Last annotation update)
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    204 AA
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01-NOV-1996 (TrEMBLrel, 01, Last sequ
01-DEC-2001 (TrEMBLrel, 19, Last anno
                           01-JUL-1997 (TrEMBLrel. 04, Created) 01-JUL-1997 (TrEMBLrel. 04, Last seq 01-JUL-2001 (TrEMBLrel. 17, Last ann Glucagon precursor. Heloderma suspectum (Gila monster).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRT;
    PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23553 MW;
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PRELIMINARY;
012955;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  204 AA;
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es 15; Conserv
                                                                                                                                  NCBI_TaxID=8554;
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NCBI_TaxID=9031;
[1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Proglucagon
                                                                                                                       Heloderma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PEPTIDE
VARSPLIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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PEPTIDE
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Q91410;
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Query Match
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Q9DDE6
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SEQUENCE FROM N.A.
MEDLINE-95295739; PubMed-7776976;
MIDLINE-95295739; PubMed-7776976;
MITOUT and chicken proglucagon: alternative splicing generates mRNA transcripts encoding glucagon-like peptide 2.";
Mol. Endocrinol. 9:267-277(1995).
EMBL; S78477; ABR34506.1; --
HSSP: PO1274; IGCN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Canis familiaris (Dog).
Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis.
NCBL_TaxID-9615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Irwin D.M.; "cDNA clouing of proglucagon from the stomach and pancreas of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TREMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Glucagon II precursor [Contains: Glucagon; glucagon-like peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Petrómyzon marinus (Sea lamprey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Hyperoartia;
Petrómyzontifórmes; Petrómyzontidae; Petrómyzon.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 41.3%; Score 86; DB 13; Length 206; Best Local Similarity 50.0%; Pred. No. 0.0016; Matches 14; Conservative 9; Mismatches 5; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF308439; AAL09425.1; -.
InterPro: IPR000332; Glucagon.
Pfam; PF00113; hormone2; 3
PROSTE; PS00260; GLUCAGON; UNKNOWN.3.
SEQUENCE 180 AA; 21114 MW; 80F6694IAFC324FD CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                          206 AA; 23875 MW; AB299E1B02FC6AA4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-2001 (TrEWBLrel. 19, Created)
01-DEC-2001 (TrEWBLrel. 19, Last sequence update)
01-MAR-2002 (TrEWBLrel. 20, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00260; GLUCAGON; 3.
SEQUENCE 206 AA; 23875 MW;
                                                                                                                                                                                                                                                                 Interpro; IPR000532; Glucagon.
Pfam; PF00123; hormone2; 3.
PRINTS; PR00275; GLUCAGON.
SMART; SM00070; GLUCA; 3.
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Q9PURO;
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Q95LG0
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MEDLINE-20022986; PubMed-10555286;
A Irvin D.M., Huner O., Youson J.H.;
A Irvin D.M., Huner O., Youson J.H.;
"Lamprey proglucagon and the Oxigin of glucagon-like peptides.";
I. Side of the Edge of t
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Eukaryota; Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi;
extinopterygii: Neopterygii: Teleostei; Ostariophysi; Cypriniformes;
Cyprinidae; Danio.
NCBL_TaxID-7955;
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MEDIINE-99425190; PubMed-10495291;
Argenton F., Zecchin E., Bortolussi M.;
Early apperance of pancreatic hormone-expressing cells in the zebrafish embryo.";
Mech. Dev. 87:217-221(1999).
ESMBL, AJ133697; GGC20108.1;
ESSP; P01274; IGCN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CHAIN 88 121 GLUCAGON-LIKE PEPTIDE 1. SEQUENCE 121 AA; 13537 MW; A85385F690DA180F CRC64;
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2011 (TrEMBLrel. 19, Last annotation update)
Glucagon polyprotein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121 AA.
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InterPro; IPR000532; Glucagon.
Pfan; PF00123; hormone2; 2.
PRINTS; PR00275; GLUCAGON.
SMART; SM0070; GLUCA; 2.
PROSITE; PS000260; GLUCAGON; 1.
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Matches 13; Conservative
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Barloy-Hubler F., Bowser L., Capela D., Galibert F., Gouzy J.,
Gurjal M., Hong A., Huizar L., Hyman R.W., Kahn D., Kahn M.L.,
Kalman S., Keating D.H., Palm C., Peck M.C., Surzycki R., Wells D.H.,
Yeh K.-C., Davis R.W., Federspiel N.A., Long S.R.,
"Nucleotide sequence and predicted functions of the entire
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei; Caranthomorpha; Acanthopterygii; Percomorpha; Perciformes; Percoidei; NCBI_TaxID=109273;
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01-DEC-2001 (TIENBLrel. 19, Created)
01-DEC-2001 (TIENBLrel. 19, Last sequence update)
01-MAR-2002 (TIENBLrel. 20, Last annotation update)
01-MAR-2002 (TIENBLrel. 20, Last annotation update)
11-MAR-2002 (TIENBLrel. 20, Last annotation update)
127 ON SMA2063.
127 ON SMA2063.
128 Plasmid pSymA (megaplasmid 1).
128 Bacteria; Protecobacteria; alpha subdivision; Rhizobiaceae group;
120 MCBL_TaxID=382;
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Pred. No. 0.0059;
5; Mismatches 15; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Al-Wahrouki A.A., Irwin D.M., Youson J.H.;
"Rock Bass Proglucagon.";
Submitted (SEP-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; Ari90499; AAG16778.1;
HSSP; P01274; 1GCN.
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706 MW; 2BDEF2867AD0475C CRC64;
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GLUCAGON-LIKE PEPTIDE 1.
GLUCAGON-LIKE PEPTIDE 2.
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                                                                                                                            01-MAR-2001 (TIEMBLrel. 16, Created)
01-MAR-2001 (TIEMBLrel. 16, Last sequence update)
01-DEC-2001 (TIEMBLrel. 19, Last annotation update)
Proglicagon (Fragment).
Ambioplites rupestris.
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Proc. Natl. Acad. Scl. U.S.A. 98:9883-9888(2001).
EMBL; AR07298; AR65785.1;
PREFPCO: IRRODIO05; Myb.DNA.binding.
PROSITE; PSOU037; MYB_1; UNKNOWN_1.
                                                                 96 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                InterPro; IPR000332; Glucagon.
Pfam; PF00123; bormone2; 2.
PRINTS; PR00275; GLUCAGON.
SMART; SR00070; GLUCA; 2.
PROSITE; PS00260; GLUCAGON; UNKNOWN_1.
                                                                     PRT;
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milarity 42.9%;
Conservative 5;
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96 AA; 11225 MW;
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                                                                     PRELIMINARY;
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Best Local Similarity
Matches 15; Conserv
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SEQUENCE 3
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SEQUENCE
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Q92XW3
RESULT 14
Q9DG43
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```

```
        Query Match
        29.8%;
        Score 62;
        DB 16;
        Length 347;

        Best Local Similarity 43.3%;
        Pred. No. 4.9;
        Matches 13;
        Conservative 5;
        Mismatches 12;
        Indels 0;
        Gaps 0;

        Qy
        10 LSYQMEERAVELEIWINGGPSSGAPPPS 39
        1
        | | | | | | | | |
        |
        Db 272 LQWSVSEBADLLSLEWVESGGPSVSEPSRS 301
```

Search completed: March 28, 2003, 08:39:25 Job time: 87 secs